

US EPA ARCHIVE DOCUMENT

**DATA EVALUATION RECORD
VEGETATIVE VIGOR EC₂₅ TEST
§123-1(b) (TIER II)**

3/24/04

3. CITATION:

Author: M. T. Christ and J. Abedi

Title: Effect on Vegetative Vigor of Non-Target Terrestrial Plants (Tier II), AE F130060 + AE F107892, Water Dispersible Granule (75.3% w/w), Including a Representative Adjuvant

Study Completion Date: August 15, 2002

Laboratory: Aventis CropScience, Ecotoxicology Department
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Research Triangle Park, NC 27709, USA

Sponsor: Aventis CropScience, Ecotoxicology Department
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Research Triangle Park, NC 27709, USA

Laboratory Report ID: CK99W504

MRID No.: 45745701

DP Barcode: D295614

- 4. REVIEWED BY:** Rebecca Bryan, Staff Scientist, Dynamac Corporation

Signature:

Date: 11/7/03

APPROVED BY: Teri Myers, Ph.D., Staff Scientist, Dynamac Corporation

Signature:

Date: 11/7/03

- 5. APPROVED BY:** Michael Davy OPP/EEED/ERBU

Signature:

Michael Dany

Date: 3/24/04



2013018

**DATA EVALUATION RECORD
VEGETATIVE VIGOR EC₂₅ TEST
§123-1(b) (TIER II)**

1. CHEMICAL: Mesosulfuron-methyl

PC Code No.: 122009

2. TEST MATERIAL: AE F130060 (a.i.) +
AE F107892 (adjuvant)

Purity: 74.73% +
3.54%

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Signature: *Rebecca Bryan*

Date: 11/7/03

APPROVED BY: Teri Myers, Ph.D., Staff Scientist, Dynamac Corporation

Signature: *Teri S. Myers*

Date: 11/7/03

5. APPROVED BY: Leo Lasota, OPP/EFED/ERBIII

Signature:

Date:

2

6. STUDY PARAMETERS:

Scientific Name of Test Organism: Dicots: *Brassica oleracea*, *Cucumis sativus*,
Lactuca sativa, *Raphanus sativus*, *Glycine max*
Lycopersicon esculentum
Monocots: *Zea mays*, *Avena sativa*, *Allium cepa*,
Triticum aestivum

Definitive Study Duration: 14 days

Type of Concentrations: Nominal

7. CONCLUSIONS:

Vegetative vigor was studied on 10 plant species after application of AE F130060 + AE F107892 75.3% (a.i. Mesosulfuron-methyl, 74.73%) at 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha. Test species included cabbage, corn, cucumber, lettuce, oat, onion, radish, soybean, tomato, and wheat. With the exception of wheat shoot length, all species and endpoints showed sensitivity to treatment, exhibiting reductions exceeding 25%. Tomato, a dicot, was the most sensitive species (based on dry weight), with an EC₂₅ of 0.18 g a.i./ha; the EC₀₅ and NOEC were 0.041 and 0.56 g a.i./ha. Corn was the most sensitive monocot (based on dry weight), with EC₂₅ of 0.47; the EC₀₅ and NOEC were 0.15 and 0.56 g a.i./ha.

This study is classified as Supplemental. This study is scientifically sound but does not fulfill the guideline requirements for vegetative vigor studies (Subdivision J, §123-1 (TIER II)).

Most sensitive monocot: Corn

Most sensitive parameter: Dry weight

NOEC: 0.56 g a.i./ha

EC₀₅: 0.15 g a.i./ha 95% C.I.: 0.059-0.39 g a.i./ha

EC₂₅: 0.47 g a.i./ha 95% C.I.: 0.27-0.84 g a.i./ha

Slope: 1.96±0.337

Most sensitive dicot: Tomato

Most sensitive parameter: Dry weight

NOEC: 0.56 g a.i./ha

EC₀₅: 0.0041 g a.i./ha 95% C.I.: 0.013-0.13 g a.i./ha

EC₂₅: 0.18 g a.i./ha 95% C.I.: 0.082-0.39 g a.i./ha

Slope: 1.52±0.229

8. ADEQUACY OF THE STUDY:

A. Classification: Supplemental

B. Rationale: This study is scientifically sound and but does not fulfill the guideline requirements for vegetative vigor studies (Subdivision J, §123-1 (TIER II)). Please see guideline deviations for explanation.

C. Repairability: None

9. GUIDELINE DEVIATIONS:

1. The herbicide being tested has an ALS inhibitor mode of action in which morphological symptoms of herbicide injury usually is seen in sensitive plants about 2 weeks after exposure to the herbicide. This delayed symptom is sometimes called the "slow death" syndrome. This study was conducted for 2 weeks. The EFED has recommended that studies that uses an ALS inhibitor herbicide br conducted for at least 3 weeks. Because this study was not conducted for sufficient amount of time, the EC25 values in this study are considered to be under reported for phytotoxic sensitivity.
2. The formulation tested in this study is not the formulation that will used in the U.S. The EFED has stated that for terrestrial plant studies, the study must use a formulation that has the highest percentage active ingredient

10. SUBMISSION PURPOSE: This study was submitted to provide data on the phytotoxicity to non-target crop species of AE F130060 + AE F107892 (a.i. mesosulfuron-methyl) after post-emergent application for the purpose of chemical registration.

11. MATERIALS AND METHODS:**A. Test Organisms**

| Guideline Criteria | Reported Information |
|--|--|
| Species: 6 dicots in 4 families, including soybean and a rootcrop; 4 monocots in 2 families, including corn. | <u>Dicots</u> : cabbage, cucumber, lettuce, radish, soybean, and tomato <u>Monocots</u> : corn, oat, onion, and wheat |
| Number of plants per repetition: | 3 plants per replicate pot (10 replicates per control and treatment group) |
| Source of seed and historical % germination of seed: | See Table 1, p. 28 for seed source information and historical % germination of seed (85-100%). |

B. Test System

| Guideline Criteria | Reported Information |
|-------------------------------------|--|
| Solvent: | N/A (adjuvant contained Synperonic™ and the safener AE F107892 at a concentration of 24 g/L) |
| Site of test: | Greenhouse #2 Tests were performed at Aventis CropScience Research Center, Ecotoxicology Laboratory, Research Triangle Park, NC. |
| Planting method/type of pot: | Plastic pots with 6 inch diameter. Corn, cucumber, ryegrass, soybean, and wheat were planted at 2.5 cm depth. Cabbage, lettuce, onion, radish, and tomato were planted at 1.3 cm depth. The sandy loam soil used for planting was a mixture of natural topsoil and sand (pH 5.7 and 1.1% organic matter). The soil was fertilized periodically throughout the study. |
| Method of application: | Track sprayer enclosed in a fume hood. |

| Guideline Criteria | Reported Information |
|-------------------------------------|---|
| Method of watering: | The pots were subirrigated an average of twice a day. The water used was tap water. |
| Growth stage at application: | Seedlings (1-5 leaf stage, p. 15) |

C. Test Design

| Guideline Criteria | Reported Information |
|--|---|
| Dose range: 2x or 3x | 3x |
| Doses: At least 5 | 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha |
| Controls: Negative and solvent | Negative control (deionized water) |
| Replicates per dose: At least 3 | 3 replicates |
| Test duration: 21 days due to ALS inhibitor herbicide | 14 days |
| Were observations made at least weekly? | Yes |
| Maximum dosage rate: | The highest treatment was equivalent to a field application rate of 15 g a.i./ha. The application rate for the spray solution was 93.5 L/ha |

12. REPORTED RESULTS:

| Guideline Criteria | Reported Information |
|---|--|
| Quality assurance and GLP compliance statements were included in the report? | Yes |
| Was a NOEC observed for each species? | Yes |
| Phytotoxic observations: | See p. 21 and Table 6, p. 33 for phytotoxicity rating system. |
| Were initial chemical concentrations measured? (Optional) | Yes, the spray trial concentrations were measured (101-106% of nominal). |

| Guideline Criteria | Reported Information |
|----------------------------------|-------------------------------|
| Were adequate raw data included? | Replicate data were provided. |

Results for the most sensitive parameter of each species**Results Synopsis****Vegetative Vigor**

| Crop | Day 21 Survival* | | Shoot length* | | Dry weight* | | Most sensitive parameter |
|----------|------------------|------------------|---------------|------------------|-------------|------------------|--------------------------|
| | NOEC | EC ₂₅ | NOEC | EC ₂₅ | NOEC | EC ₂₅ | |
| Cabbage | 15 | ND | 0.56 | 1.77 | 0.56 | 0.42 | Dry weight |
| Corn | 1.67 | ND | 0.56 | 1.08 | 0.56 | 0.62 | Dry weight |
| Cucumber | 5 | ND | 0.19 | 1.76 | 0.19 | 0.67 | Dry weight |
| Lettuce | 15 | ND | 0.19 | 0.41 | 0.19 | 0.22 | Dry weight |
| Oat | 5 | ND | 0.56 | 1.32 | 0.56 | 0.80 | Dry weight |
| Onion | 5 | ND | 1.67 | 3.39 | 1.67 | 2.11 | Dry weight |
| Radish | 1.67 | ND | 0.19 | 0.51 | 0.19 | 0.36 | Dry weight |
| Soybean | 15 | ND | 0.19 | 0.79 | 0.19 | 0.24 | Dry weight |
| Tomato | 1.67 | ND | 0.19 | 0.47 | 0.19 | 0.31 | Dry weight |
| Wheat | 15 | ND | 5 | >15 | 5 | 9.2 | Dry weight |

ND = Not determined.

* Units are g a.i./ha

Morphological Observations (negative percent reductions indicate promoted growth)

Cabbage: Seedling survival was 100% for all treatment groups by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were -4, 0, -1, 26, 42, and 47% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were -14, -9, 6, 65, 75, and 84% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in

the 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the phytotoxic effects including chlorosis, leaf cupping, and height inhibition were significant in the 1.67, 5, and 15 g a.i./ha treatment groups.

Corn: Seedling survival was 100, 100, 100, 100, 40, and 7% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 2, 15, 9, 54, 81, and 92% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 1, 22, 22, 67, 91, and 92% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included leaf cupping in the 1.67 g a.i./ha treatment group, height inhibition in the 1.67, 5, and 15 g a.i./ha treatment groups, and necrosis in the 5 and 15 g a.i./ha treatment groups.

Cucumber: Seedling survival was 100, 100, 100, 100, 97, and 63% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 5, 2, 14, 27, 36, and 35% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 18, 3, 24, 47, 60, and 62% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis in the 1.67 g a.i./ha treatment group, leaf cupping and height inhibition in the 1.67, 5, and 15 g a.i./ha treatment groups, and necrosis in the 15 g a.i./ha treatment groups.

Lettuce: Seedling survival was 100% for all treatment groups by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 5, 1, 17, 52, 67, and 68% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 12, 6, 35, 70, 80, and 86% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in

the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis and height inhibition in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, and necrosis in the 1.67, 5, and 15 g a.i./ha treatment groups.

Oat: Seedling survival was 100, 100, 100, 100, 87, and 50% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 6, -2, 1, 37, 61, and 69% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 3, -9, -1, 59, 82, and 88% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis and leaf twisting in the 1.67 g a.i./ha treatment group, height inhibition in the 1.67, 5, and 15 g a.i./ha treatment groups, and necrosis in the 5 and 15 g a.i./ha treatment groups.

Onion: Seedling survival was 100, 100, 100, 100, 93, and 63% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 0, 7, 3, 10, 39, and 47% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 1, 5, -7, 15, 54, and 69% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 5 and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included leaf malformation and height inhibition in the 5 and 15 g a.i./ha treatment groups, and necrosis in the 15 g a.i./ha treatment group.

Radish: Seedling survival was 100, 100, 87, 97, 33, and 3% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were -5, -10, 30, 48, 60, and 75% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were -5, -6,

44, 57, 57, and 75% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis and leaf malformation in the 0.56 and 1.67 g a.i./ha treatment groups, and height inhibition and necrosis in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

Soybean: Seedling survival was 100% for all treatment groups by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were -1, 0, 19, 39, 43, and 45% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 1, 1, 27, 65, 77, and 74% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis, leaf malformation, height inhibition, and necrosis in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

Tomato: Seedling survival was 100, 100, 100, 100, 43, and 7% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were 8, 3, 34, 61, 77, and 78% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were 13, 5, 55, 86, 92, and 90% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups.

By 14 days, the significant phytotoxic effects included chlorosis and leaf curling in the 0.56 and 1.67 g a.i./ha treatment groups, height inhibition in the 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, and necrosis in the 1.67, 5, and 15 g a.i./ha treatment groups.

Wheat: Seedling survival was 100% for all treatment groups by 14 days; control survival was 100%.

By 14 days, the percent reductions of mean plant heights were -12, -6, -5, 0, 6, and 19% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The percent reductions of mean dry weights were -11, -23, -34, -12,

-3, and 33% in the 0.06, 0.19, 0.56, 1.67, 5, and 15 g a.i./ha treatment groups, respectively, compared to the control group. The height and dry weight inhibitions were significant in the 15 g a.i./ha treatment group.

By 14 days, the significant phytotoxic effects included leaf curling and height inhibition in the 15 g a.i./ha treatment group.

Statistical Results

Statistical Method: The percent inhibition was calculated using the equation on p. 22. The survival data was analyzed using the Fisher's Exact Test. The Shapiro-Wilk's test was used to determine normal distribution and the Bartlett's Test to determine homogeneity of variance for all data. The treatment group data was compared to the control using the Dunnett's one way analysis of variance. If transformations were not successful, the data were analyzed using Steele's Rank Sum Test. The EC₂₅ and EC₅₀ were calculated using non-linear regression based on Bruce and Versteeg or a linear interpolation program. The SAS® Procedure NLIN (Version 6.12) computer software was used to perform statistical analyses. The linear interpolation analyses were performed using Inhibition Concentration Approach (ICp)(Version 2.0).

Most sensitive monocot: Corn

Most sensitive parameter: Dry weight

NOEC: 0.56 g a.i./ha

EC₂₅: 0.62 g a.i./ha 95% C.I.: 0.35-0.90 g a.i./ha

EC₅₀: 1.17 g a.i./ha 95% C.I.: 0.86-1.61 g a.i./ha

Slope: Not reported

Most sensitive dicot: Lettuce

Most sensitive parameter: Dry weight

NOEC: 0.19 g a.i./ha

EC₂₅: 0.22 g a.i./ha 95% C.I.: 0.09-0.55 g a.i./ha

EC₅₀: 0.92 g a.i./ha 95% C.I.: 0.50-1.70 g a.i./ha

Slope: Not reported

13. REVIEWER'S VERIFICATION OF STATISTICAL RESULTS:

Statistical Method: Shoot length and dry weight data were analyzed to determine if they satisfied the assumptions of ANOVA (i.e., normal distribution and homogeneity of variances). If they did, the NOEC was determined using ANOVA, followed by Bonferroni's t-test (unequal replicates, non-monotonic response), Dunnett's test (equal replicates, non-monotonic response), or William's test (monotonic response). If the data

did not meet these assumptions, transformations (e.g., square-root, inverse square-root, or natural log) were attempted. If these transformations were successful, the NOEC was determined using a method described above. If the transformations were not successful, the NOEC was determined using the non-parametric Kruskal-Wallis test. These analyses were conducted using TOXSTAT statistical software. The EC₀₅ and EC₂₅ values and their 95% confidence intervals and slopes were determined using the Probit method via Nuthatch statistical software.

Morphological Symptoms NOAEC - 14 DAT (Results in gm ai/ha)

Cabbage - 0.06,

Corn - ND since lowest dose has height inhibition

Cucumber - 0.19

Lettuce - 0.19

Oat - 1.67

Onion - 0.06

Radish - ND since at lowest dose has necrosis
chlorosis and leaf malformation

Soybean - 0.19

Tomato - 0.06

Wheat - 5.0

Results synopsis

| Crop | Shoot length* | | | Dry weight* | | | Most sensitive parameter |
|----------|-------------------|------------------|-------------------|-------------------|------------------|-------------------|--------------------------|
| | NOEC | EC ₀₅ | EC ₂₅ | NOEC | EC ₀₅ | EC ₂₅ | |
| Cabbage | 0.56 | 0.12 | 1.8 | 0.56 | 0.072 | 0.42 | Dry weight |
| Corn | 0.06 ^a | 0.27 | 0.82 ^a | 0.56 | 0.15 | 0.47 ^a | Dry weight |
| Cucumber | 0.19 | 0.028 | 1.9 ^b | 0.19 | 0.025 | 0.46 ^a | Dry weight |
| Lettuce | 0.19 | 0.033 | 0.41 | 0.19 | 0.028 | 0.22 | Dry weight |
| Oat | 0.56 | 0.16 | 1.0 ^a | 0.56 | 0.16 | 0.63 ^a | Dry weight |
| Onion | 1.67 | 0.40 | 3.4 | 1.67 | 0.50 | 2.1 | Dry weight |
| Radish | 0.19 | 0.074 | 0.51 | 0.19 | 0.028 | 0.29 ^a | Dry weight |
| Soybean | 0.19 | 0.019 | 0.79 | 0.19 | 0.021 | 0.24 | Dry weight |
| Tomato | 0.19 | 0.044 | 0.31 ^a | 0.56 ^b | 0.041 | 0.18 ^a | Dry weight |
| Wheat | 5 | 1.9 | >15 | 5 | 2.9 | 8.7 ^a | Dry weight |

^aThe reviewer's estimate was lower than the study authors'.

^bThe reviewer's estimate was higher than the study authors'.

*units are g a.i./ha

EC_x values, confidence intervals, and slopes

| Species | Shoot length* | | | | | Dry weight* | | | | |
|----------|------------------|---------------------|-------------------|---------------------|--------------|------------------|---------------------|-------------------|---------------------|--------------|
| | EC ₀₅ | Confidence interval | EC ₂₅ | Confidence interval | Slope | EC ₀₅ | Confidence interval | EC ₂₅ | Confidence interval | Slope |
| Cabbage | 0.12 | 0.038-0.40 | 1.8 | 1.0-3.1 | 0.840±0.104 | 0.072 | 0.029-0.17 | 0.42 | 0.24-0.74 | 1.26±0.123 |
| Corn | 0.27 | 0.15-0.47 | 0.82 ^a | 0.59-1.1 | 2.00±0.234 | 0.15 | 0.059-0.39 | 0.47 ^a | 0.27-0.84 | 1.96±0.337 |
| Cucumber | 0.028 | 0.0039-0.21 | 1.9 ^b | 0.83-4.4 | 0.531±0.0844 | 0.025 | 0.0029-0.22 | 0.46 ^a | 0.14-1.5 | 0.765±0.137 |
| Lettuce | 0.033 | 0.0094-0.12 | 0.41 | 0.20-0.87 | 0.883±0.0957 | 0.028 | 0.0067-0.11 | 0.22 | 0.087-0.55 | 1.08±0.139 |
| Oat | 0.16 | 0.078-0.35 | 1.0 ^b | 0.68-1.6 | 1.21±0.116 | 0.16 | 0.074-0.35 | 0.63 ^a | 0.39-1.0 | 1.63±0.182 |
| Onion | 0.40 | 0.10-1.6 | 3.4 | 1.9-6.2 | 1.05±0.208 | 0.50 | 0.10-2.5 | 2.1 | 0.86-5.2 | 1.56±0.422 |
| Radish | 0.074 | 0.023-0.24 | 0.51 | 0.27-0.97 | 1.16±0.177 | 0.028 | 0.0045-0.18 | 0.29 ^a | 0.10-0.81 | 0.962±0.187 |
| Soybean | 0.019 | 0.0043-0.082 | 0.79 | 0.38-1.6 | 0.598±0.0655 | 0.021 | 0.0059-0.077 | 0.24 | 0.11-0.53 | 0.928±0.0995 |
| Tomato | 0.044 | 0.019-0.10 | 0.31 ^a | 0.19-0.51 | 1.15±0.107 | 0.041 | 0.013-0.13 | 0.18 ^a | 0.082-0.39 | 1.52±0.229 |
| Wheat | 1.9 | 0.63-6.0 | >15 | N/A | 1.05±0.29 | 2.9 | 0.74-11 | 8.7 ^a | 5.2-15 | 2.01±0.840 |

^aThe reviewer's estimate was lower than the study authors'.^bThe reviewer's estimate was higher than the study authors'.

*units are g a.i./ha

Most sensitive monocot: Corn

Most sensitive parameter: Dry weight

NOEC: 0.56 g a.i./ha

EC₀₅: 0.15 g a.i./ha 95% C.I.: 0.059-0.39 g a.i./haEC₂₅: 0.47 g a.i./ha 95% C.I.: 0.27-0.84 g a.i./ha

Slope: 1.96±0.337

Most sensitive dicot: Tomato

Most sensitive parameter: Dry weight

NOEC: 0.56 g a.i./ha

EC₀₅: 0.041 g a.i./ha 95% C.I.: 0.013-0.13 g a.i./haEC₂₅: 0.18 g a.i./ha 95% C.I.: 0.082-0.39 g a.i./ha

Slope: 1.52±0.229

14. REVIEWER'S COMMENTS:

The herbicide being tested has an ALS inhibitor mode of action in which morphological symptoms of herbicide injury usually is seen in sensitive plants about

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2 weeks after exposure to the herbicide. This delayed symptom is sometimes called the "slow death" syndrome. This study was conducted for 2 weeks. The EFED has recommended that studies that uses an ALS inhibitor herbicide be conducted for at least 3 weeks. Because this study was not conducted for sufficient amount of time, the EC₂₅ values in this study are considered to be under reported for phytotoxic sensitivity.

The formulation tested in this study is not the formulation that will be used in the U.S. The EFED has stated that for terrestrial plant studies, the study must use a formulation that has the highest percentage active ingredient

The reviewer's conclusions regarding the most sensitive species (tomato, a dicot) differed from the study authors' (lettuce); furthermore, the reviewer's EC₂₅ estimate for corn (the most sensitive monocot) was slightly lower than the study authors' estimate. Differences between the reviewer's and the study authors' estimates can be attributed to the different statistical methods which were used to derive these estimates. Because the reviewer's analysis provided EC₀₅ values and slopes for all estimates, the reviewer's values are reported in the Conclusions section.

The morphological symptoms NOAEC differ from those NOAEC found for cabbage, onion and tomato. Each of these species have the NOAEC of 0.06 gm ai/ha which is more sensitive than the NOAEC found for measured length and weights.

The definitive study for cabbage, corn, oat, radish, and wheat was conducted from March 26, 2002 to April 9, 2002. The definitive study for cucumber, lettuce, tomato, and soybean was conducted from March 28, 2002 to April 11, 2002. The temperature ranged from 21.2 to 26.5°C and the humidity ranged from 21.6 to 67.5%. The light intensity was 157.7 to 453.9 $\mu\text{mol m}^{-2}\text{s}^{-1}$.

The definitive study for onion was conducted from June 4, 2002 to June 18, 2002. The temperature ranged from 26.5 to 29.3°C and the humidity ranged from 36.8 to 60.8%. The light intensity was 182.9 to 462.8 $\mu\text{mol m}^{-2}\text{s}^{-1}$.

The oat dry weight means were incorrectly reported in Table 10, p. 66 (values reported on this page are the means for onion dry weight).

A non-GLP range finding test and other previous definitive trials were conducted to determine the treatment concentrations for the final definitive study. The previous trials were not reported due to low analytical recoveries and a change in the adjuvant/safener rate.

This study was conducted in accordance with USEPA Good Laboratory Practice

DP Barcode: D295614

MRID No.: 45745701

Standards with the exception of the collection of data for the water and soil contaminant and screening analyses (p. 3). The study included a Quality Assurance statement.

15. REFERENCES:

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APPENDIX I. OUTPUT FROM REVIEWER'S STATISTICAL VERIFICATION:

cabbage length

File: 5701bl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|----------|---------|--------|
| Between | 6 | 921.288 | 153.548 | 62.090 |
| Within (Error) | 63 | 155.810 | 2.473 | |
| Total | 69 | 1077.098 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho: All groups equal

cabbage length

File: 5701bl Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|----------|-----|
| 1 | control | 17.620 | 17.620 | | |
| 2 | 0.06 | 18.380 | 18.380 | -1.081 | |
| 3 | 0.19 | 17.590 | 17.590 | 0.043 | |
| 4 | 0.56 | 17.780 | 17.780 | -0.228 | |
| 5 | 1.67 | 12.960 | 12.960 | 6.626 * | * |
| 6 | 5 | 10.210 | 10.210 | 10.536 * | * |
| 7 | 15 | 9.300 | 9.300 | 11.830 * | * |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

cabbage length

File: 5701bl Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 1.653 | 9.4 | -0.760 |
| 3 | 0.19 | 10 | 1.653 | 9.4 | 0.030 |
| 4 | 0.56 | 10 | 1.653 | 9.4 | -0.160 |
| 5 | 1.67 | 10 | 1.653 | 9.4 | 4.660 |
| 6 | 5 | 10 | 1.653 | 9.4 | 7.410 |
| 7 | 15 | 10 | 1.653 | 9.4 | 8.320 |

cabbage length

File: 5701bl Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 17.620 | 17.620 | 18.000 |
| 2 | 0.06 | 10 | 18.380 | 18.380 | 18.000 |
| 3 | 0.19 | 10 | 17.590 | 17.590 | 17.685 |
| 4 | 0.56 | 10 | 17.780 | 17.780 | 17.685 |
| 5 | 1.67 | 10 | 12.960 | 12.960 | 12.960 |
| 6 | 5 | 10 | 10.210 | 10.210 | 10.210 |
| 7 | | 15 | 9.300 | 9.300 | 9.300 |

cabbage length

File: 5701bl

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 18.000 | | | | |
| 0.06 | 18.000 | 0.540 | | 1.67 | k= 1, v=63 |
| 0.19 | 17.685 | 0.092 | | 1.75 | k= 2, v=63 |
| 0.56 | 17.685 | 0.092 | | 1.77 | k= 3, v=63 |
| 1.67 | 12.960 | 6.626 | * | 1.78 | k= 4, v=63 |
| 5 | 10.210 | 10.536 | * | 1.79 | k= 5, v=63 |
| 15 | 9.300 | 11.830 | * | 1.79 | k= 6, v=63 |

s = 1.573

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.12 | 0.038 | 0.40 | 0.26 | 0.31 |
| EC10 | 0.34 | 0.13 | 0.86 | 0.20 | 0.39 |
| EC25 | 1.8 | 1.0 | 3.1 | 0.12 | 0.57 |
| EC50 | 11. | 8.2 | 15. | 0.069 | 0.73 |

Slope = 0.840 Std.Err. = 0.104

!!!Poor fit: p < 0.001 based on DF= 4.00 63.0

5701BL : cabbage length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|-----------|------------|-------------|----------------|---------|
| 0.00 | 10.0 | 17.6 | 18.5 | -0.925 | 100. | 0.00 |
| 0.0600 | 10.0 | 18.4 | 18.0 | 0.356 | 97.2 | 2.81 |
| 0.190 | 10.0 | 17.6 | 17.3 | 0.310 | 93.2 | 6.82 |
| 0.560 | 10.0 | 17.8 | 16.0 | 1.77 | 86.3 | 13.7 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | | |
|------|------|------|------|-------|------|------|
| 1.67 | 10.0 | 13.0 | 14.0 | -1.08 | 75.7 | 24.3 |
| 5.00 | 10.0 | 10.2 | 11.4 | -1.23 | 61.7 | 38.3 |
| 15.0 | 10.0 | 9.30 | 8.51 | 0.794 | 45.9 | 54.1 |

cabbage dry weight

File: 5701bw Transform: SQUARE ROOT(Y)

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|--------|-------|---------|
| Between | 6 | 13.075 | 2.179 | 136.188 |
| Within (Error) | 63 | 0.982 | 0.016 | |
| Total | 69 | 14.057 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

cabbage dry weight

File: 5701bw Transform: SQUARE ROOT(Y)

DUNNETTS TEST - TABLE 1 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED | | MEAN CALCULATED IN | | T STAT | SIG |
|-------|----------------|-------------|----------------|--------------------|--|--------|-----|
| | | MEAN | ORIGINAL UNITS | | | | |
| 1 | control | 1.621 | | 2.653 | | | |
| 2 | 0.06 | 1.738 | | 3.031 | | -2.079 | |
| 3 | 0.19 | 1.694 | | 2.887 | | -1.300 | |
| 4 | 0.56 | 1.575 | | 2.493 | | 0.799 | |
| 5 | 1.67 | 0.955 | | 0.930 | | 11.762 | * |
| 6 | 5 | 0.810 | | 0.667 | | 14.331 | * |
| 7 | 15 | 0.650 | | 0.429 | | 17.164 | * |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

cabbage dry weight

File: 5701bw Transform: SQUARE ROOT(Y)

DUNNETTS TEST - TABLE 2 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of | DIFFERENCE |
|-------|----------------|----------------|--------------------------------------|---------|--------------|
| | | | | CONTROL | FROM CONTROL |
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.413 | 15.6 | -0.378 |
| 3 | 0.19 | 10 | 0.413 | 15.6 | -0.234 |
| 4 | 0.56 | 10 | 0.413 | 15.6 | 0.160 |
| 5 | 1.67 | 10 | 0.413 | 15.6 | 1.723 |
| 6 | 5 | 10 | 0.413 | 15.6 | 1.986 |
| 7 | 15 | 10 | 0.413 | 15.6 | 2.224 |

cabbage dry weight
 File: 5701bw Transform: SQUARE ROOT(Y)

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 2.653 | 1.621 | 1.684 |
| 2 | 0.06 | 10 | 3.031 | 1.738 | 1.684 |
| 3 | 0.19 | 10 | 2.887 | 1.694 | 1.684 |
| 4 | 0.56 | 10 | 2.493 | 1.575 | 1.575 |
| 5 | 1.67 | 10 | 0.930 | 0.955 | 0.955 |
| 6 | 5 | 10 | 0.667 | 0.810 | 0.810 |
| 7 | | 15 | 0.429 | 0.650 | 0.650 |

cabbage dry weight
 File: 5701bw Transform: SQUARE ROOT(Y)

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 1.684 | | | | |
| 0.06 | 1.684 | 1.141 | | 1.67 | k= 1, v=63 |
| 0.19 | 1.684 | 1.141 | | 1.75 | k= 2, v=63 |
| 0.56 | 1.575 | 0.809 | | 1.77 | k= 3, v=63 |
| 1.67 | 0.955 | 11.916 | * | 1.78 | k= 4, v=63 |
| 5 | 0.810 | 14.518 | * | 1.79 | k= 5, v=63 |
| 15 | 0.650 | 17.388 | * | 1.79 | k= 6, v=63 |

s = 0.125

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.072 | 0.029 | 0.17 | 0.19 | 0.41 |
| EC10 | 0.14 | 0.064 | 0.30 | 0.17 | 0.46 |
| EC25 | 0.42 | 0.24 | 0.74 | 0.12 | 0.56 |
| EC50 | 1.4 | 0.98 | 2.1 | 0.082 | 0.69 |

Slope = 1.26 Std.Err. = 0.123

!!!Poor fit: p < 0.001 based on DF= 4.00 63.0

5701BW : cabbage dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. - Pred. | Pred. %Control | %Change |
|------|--------|-----------|------------|--------------|----------------|---------|
| 0.00 | 10.0 | 2.65 | 3.02 | -0.368 | 100. | 0.00 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | | |
|--------|------|-------|-------|---------|------|------|
| 0.0600 | 10.0 | 3.03 | 2.90 | 0.134 | 95.9 | 4.08 |
| 0.190 | 10.0 | 2.89 | 2.62 | 0.270 | 86.6 | 13.4 |
| 0.560 | 10.0 | 2.49 | 2.10 | 0.388 | 69.7 | 30.3 |
| 1.67 | 10.0 | 0.930 | 1.41 | -0.479 | 46.6 | 53.4 |
| 5.00 | 10.0 | 0.667 | 0.743 | -0.0767 | 24.6 | 75.4 |
| 15.0 | 10.0 | 0.429 | 0.298 | 0.131 | 9.85 | 90.2 |

corn length

File: 5701nl

Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|-----------|----------|--------|
| Between | 5 | 20805.863 | 4161.173 | 49.139 |
| Within (Error) | 49 | 4149.396 | 84.682 | |
| Total | 54 | 24955.259 | | |

Critical F value = 2.45 (0.05, 5, 40)

Since F > Critical F REJECT Ho: All groups equal

corn length

File: 5701nl

Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 1 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|--------|-----|
| 1 | control | 74.030 | 74.030 | | |
| 2 | 0.06 | 72.730 | 72.730 | 0.316 | |
| 3 | 0.19 | 62.980 | 62.980 | 2.685 | * |
| 4 | 0.56 | 67.170 | 67.170 | 1.667 | |
| 5 | 1.67 | 34.310 | 34.310 | 9.652 | * |
| 6 | 5 | 14.720 | 14.720 | 11.767 | * |

Bonferroni T table value = 2.42 (1 Tailed Value, P=0.05, df=40, 5)

corn length

File: 5701nl

Transform: NO TRANSFORMATION.

BONFERRONI T-TEST - TABLE 2 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 9.976 | 13.5 | 1.300 |
| 3 | 0.19 | 10 | 9.976 | 13.5 | 11.050 |
| 4 | 0.56 | 10 | 9.976 | 13.5 | 6.860 |
| 5 | 1.67 | 10 | 9.976 | 13.5 | 39.720 |
| 6 | 5 | 5 | 12.218 | 16.5 | 59.310 |

22

Corn length
File: 5701nl Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 74.030 | 74.030 | 74.030 |
| 2 | 0.06 | 10 | 72.730 | 72.730 | 72.730 |
| 3 | 0.19 | 10 | 62.980 | 62.980 | 65.075 |
| 4 | 0.56 | 10 | 67.170 | 67.170 | 65.075 |
| 5 | 1.67 | 10 | 34.310 | 34.310 | 34.310 |
| 6 | | 5 | 14.720 | 14.720 | 14.720 |

Corn length
File: 5701nl Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 74.030 | | | | |
| 0.06 | 72.730 | 0.316 | | 1.68 | k= 1, v=49 |
| 0.19 | 65.075 | 2.176 | * | 1.76 | k= 2, v=49 |
| 0.56 | 65.075 | 2.176 | * | 1.79 | k= 3, v=49 |
| 1.67 | 34.310 | 9.652 | * | 1.80 | k= 4, v=49 |
| 5 | 14.720 | 11.767 | * | 1.80 | k= 5, v=49 |

s = 9.202

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.27 | 0.15 | 0.47 | 0.12 | 0.57 |
| EC10 | 0.41 | 0.25 | 0.65 | 0.10 | 0.63 |
| EC25 | 0.82 | 0.59 | 1.1 | 0.070 | 0.72 |
| EC50 | 1.8 | 1.5 | 2.2 | 0.043 | 0.82 |

Slope = 2.00 Std.Err. = 0.234

!!! Poor fit: p = 0.011 based on DF= 3.0 49.

5701NL : corn length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|-----------|------------|-------------|----------------|---------|
| 0.00 | 10.0 | 74.0 | 71.9 | 2.10 | 100. | 0.00 |
| 0.0600 | 10.0 | 72.7 | 71.8 | 0.918 | 99.8 | 0.163 |
| 0.190 | 10.0 | 63.0 | 70.1 | -7.07 | 97.4 | 2.61 |
| 0.560 | 10.0 | 67.2 | 60.6 | 6.59 | 84.2 | 15.8 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | | |
|------|------|------|------|-------|------|------|
| 1.67 | 10.0 | 34.3 | 37.6 | -3.24 | 52.2 | 47.8 |
| 5.00 | 5.00 | 14.7 | 13.3 | 1.42 | 18.5 | 81.5 |

corn dry weight

File: 5701nw Transform: 1/(SQUARE ROOT(Y))

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|-------|-------|--------|
| Between | 5 | 6.967 | 1.393 | 73.316 |
| Within (Error) | 49 | 0.932 | 0.019 | |
| Total | 54 | 7.899 | | |

Critical F value = 2.45 (0.05, 5, 40)

Since F > Critical F REJECT Ho: All groups equal

corn dry weight

File: 5701nw Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST**TABLE 1 OF 2**

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|---------|-----|
| 1 | control | 0.535 | 3.718 | | |
| 2 | 0.06 | 0.569 | 3.373 | -0.549 | |
| 3 | 0.19 | 0.646 | 2.801 | -1.805 | |
| 4 | 0.56 | 0.624 | 2.680 | -1.444 | |
| 5 | 1.67 | 0.974 | 1.135 | -7.115 | |
| 6 | 5 | 1.793 | 0.319 | -16.658 | |

Bonferroni T table value = 2.42 (1 Tailed Value, P=0.05, df=40, 5)

corn dry weight

File: 5701nw Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST**TABLE 2 OF 2**

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|----------------|--------------------------------------|-----------------|----------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | -3.231 | -86.9 | 0.345 |
| 3 | 0.19 | 10 | -3.231 | -86.9 | 0.917 |
| 4 | 0.56 | 10 | -3.231 | -86.9 | 1.038 |
| 5 | 1.67 | 10 | -3.231 | -86.9 | 2.583 |
| 6 | 5 | 5 | -4.575 | -123.1 | 3.399 |

corn dry weight

File: 5701nw Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|------------------|---------------------|--------------------|
| 1 | control | 10 | 3.718 | 0.535 | 0.535 |
| 2 | 0.06 | 10 | 3.373 | 0.569 | 0.569 |
| 3 | 0.19 | 10 | 2.801 | 0.646 | 0.635 |
| 4 | 0.56 | 10 | 2.680 | 0.624 | 0.635 |
| 5 | 1.67 | 10 | 1.135 | 0.974 | 0.974 |
| 6 | | 5 | 0.319 | 1.793 | 1.793 |

corn dry weight

File: 5701nw

Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|--------------------|-------------------|--------------|-------------------|-----------------------|
| control | 0.535 | | | | |
| 0.06 | 0.569 | 0.549 | | 1.68 | k= 1, v=49 |
| 0.19 | 0.635 | 1.624 | | 1.76 | k= 2, v=49 |
| 0.56 | 0.635 | 1.624 | | 1.79 | k= 3, v=49 |
| 1.67 | 0.974 | 7.113 | * | 1.80 | k= 4, v=49 |
| 5 | 1.793 | 16.654 | * | 1.80 | k= 5, v=49 |

s = 0.138

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 0.15 | 0.059 | 0.39 | 0.20 | 0.39 |
| EC10 | 0.23 | 0.10 | 0.52 | 0.17 | 0.45 |
| EC25 | 0.47 | 0.27 | 0.84 | 0.12 | 0.56 |
| EC50 | 1.0 | 0.73 | 1.5 | 0.079 | 0.69 |

Slope = 1.96 Std.Err. = 0.337

Goodness of fit: p = 0.30 based on DF= 3.0 49.

5701NW : corn dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 3.72 | 3.45 | 0.263 | 100. | 0.00 |
| 0.0600 | 10.0 | 3.37 | 3.43 | -0.0555 | 99.2 | 0.755 |
| 0.190 | 10.0 | 2.80 | 3.20 | -0.400 | 92.7 | 7.35 |
| 0.560 | 10.0 | 2.68 | 2.43 | 0.252 | 70.3 | 29.7 |
| 1.67 | 10.0 | 1.14 | 1.20 | -0.0598 | 34.6 | 65.4 |
| 5.00 | 5.00 | 0.319 | 0.318 | 0.00102 | 9.21 | 90.8 |

DP Barcode: D295614

MRID No.: 45745701

cucumber length

File: 5701cl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|---------|--------|--------|
| Between | 6 | 569.840 | 94.973 | 27.275 |
| Within (Error) | 60 | 208.932 | 3.482 | |
| Total | 66 | 778.772 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho: All groups equal

cucumber length

File: 5701cl Transform: NO TRANSFORMATION

BONFERRONI T-TEST

TABLE 1 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|--------------------------------------|---------|-----|
| 1 | control | 20.750 | 20.750 | | |
| 2 | 0.06 | 19.560 | 19.560 | 1.426 | |
| 3 | 0.19 | 20.190 | 20.190 | 0.671 | |
| 4 | 0.56 | 17.810 | 17.810 | 3.523 * | |
| 5 | 1.67 | 15.060 | 15.060 | 6.818 * | |
| 6 | 5 | 13.270 | 13.270 | 8.963 * | |
| 7 | 15 | 13.500 | 13.500 | 7.884 * | |

Bonferroni T table value = 2.46 (1 Tailed Value, P=0.05, df=60, 6)

cucumber length

File: 5701cl Transform: NO TRANSFORMATION

BONFERRONI T-TEST

TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|----------------|--------------------------------------|-----------------|----------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 2.055 | 9.9 | 1.190 |
| 3 | 0.19 | 10 | 2.055 | 9.9 | 0.560 |
| 4 | 0.56 | 10 | 2.055 | 9.9 | 2.940 |
| 5 | 1.67 | 10 | 2.055 | 9.9 | 5.690 |
| 6 | 5 | 10 | 2.055 | 9.9 | 7.480 |
| 7 | 15 | 7 | 2.265 | 10.9 | 7.250 |

cucumber length

File: 5701cl Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|------------------|---------------------|--------------------|
| 1 | control | 10 | 20.750 | 20.750 | 20.750 |
| 2 | 0.06 | 10 | 19.560 | 19.560 | 19.875 |
| 3 | 0.19 | 10 | 20.190 | 20.190 | 19.875 |
| 4 | 0.56 | 10 | 17.810 | 17.810 | 17.810 |
| 5 | 1.67 | 10 | 15.060 | 15.060 | 15.060 |
| 6 | 5 | 10 | 13.270 | 13.270 | 13.365 |
| 7 | 15 | 7 | 13.500 | 13.500 | 13.365 |

cucumber length

File: 5701cl

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|--------------------|-------------------|--------------|-------------------|-----------------------|
| control | 20.750 | | | | |
| 0.06 | 19.875 | 1.048 | | 1.67 | k= 1, v=60 |
| 0.19 | 19.875 | 1.048 | | 1.75 | k= 2, v=60 |
| 0.56 | 17.810 | 3.523 | * | 1.77 | k= 3, v=60 |
| 1.67 | 15.060 | 6.818 | * | 1.78 | k= 4, v=60 |
| 5 | 13.365 | 8.850 | * | 1.79 | k= 5, v=60 |
| 15 | 13.365 | 8.031 | * | 1.79 | k= 6, v=60 |

s = 1.866

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 0.028 | 0.0039 | 0.21 | 0.43 | 0.14 |
| EC10 | 0.14 | 0.030 | 0.64 | 0.33 | 0.22 |
| EC25 | 1.9 | 0.83 | 4.4 | 0.18 | 0.43 |
| EC50 | 36. | 17. | 73. | 0.16 | 0.49 |

Slope = 0.531 Std.Err. = 0.0844

!!!Poor fit: p = 0.0031 based on DF= 4.0 60.

5701CL : cucumber length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 20.8 | 21.2 | -0.406 | 100. | 0.00 |
| 0.0600 | 10.0 | 19.6 | 19.7 | -0.107 | 93.0 | 7.04 |
| 0.190 | 10.0 | 20.2 | 18.8 | 1.44 | 88.6 | 11.4 |
| 0.560 | 10.0 | 17.8 | 17.6 | 0.229 | 83.1 | 16.9 |
| 1.67 | 10.0 | 15.1 | 16.1 | -1.02 | 76.0 | 24.0 |
| 5.00 | 10.0 | 13.3 | 14.3 | -1.01 | 67.5 | 32.5 |

DP Barcode: D295614

MRID No.: 45745701

15.0 7.00 13.5 12.3 1.24 57.9 42.1

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

cucumber dry weight

File: 5701cw Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|--------|-------|--------|
| Between | 6 | 27.346 | 4.558 | 19.232 |
| Within (Error) | 60 | 14.194 | 0.237 | |
| Total | 66 | 41.540 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho: All groups equal

cucumber dry weight

File: 5701cw Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 1 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|---------|-----|
| 1 | control | 2.696 | 2.696 | | |
| 2 | 0.06 | 2.230 | 2.230 | 2.143 | |
| 3 | 0.19 | 2.646 | 2.646 | 0.230 | |
| 4 | 0.56 | 2.061 | 2.061 | 2.916 * | |
| 5 | 1.67 | 1.443 | 1.443 | 5.754 * | |
| 6 | 5 | 1.084 | 1.084 | 7.405 * | |
| 7 | 15 | 1.023 | 1.023 | 6.974 * | |

Bonferroni T table value = 2.46 (1 Tailed Value, P=0.05, df=60, 6)

cucumber dry weight

File: 5701cw Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 2 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.536 | 19.9 | 0.467 |
| 3 | 0.19 | 10 | 0.536 | 19.9 | 0.050 |
| 4 | 0.56 | 10 | 0.536 | 19.9 | 0.635 |
| 5 | 1.67 | 10 | 0.536 | 19.9 | 1.253 |
| 6 | 5 | 10 | 0.536 | 19.9 | 1.612 |
| 7 | 15 | 7 | 0.591 | 21.9 | 1.673 |

cucumber dry weight

File: 5701cw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL | TRANSFORMED | ISOTONIZED |
|-------|----------------|----|----------|-------------|------------|
| | | | MEAN | MEAN | MEAN |
| 1 | control | 10 | 2.696 | 2.696 | 2.696 |
| 2 | 0.06 | 10 | 2.230 | 2.230 | 2.438 |
| 3 | 0.19 | 10 | 2.646 | 2.646 | 2.438 |
| 4 | 0.56 | 10 | 2.061 | 2.061 | 2.061 |
| 5 | 1.67 | 10 | 1.443 | 1.443 | 1.443 |
| 6 | 5 | 10 | 1.084 | 1.084 | 1.084 |
| 7 | 15 | 7 | 1.023 | 1.023 | 1.023 |

cucumber dry weight

File: 5701cw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|--------------------|-------------------|--------------|-------------------|-----------------------|
| control | 2.696 | | | | |
| 0.06 | 2.438 | 1.188 | | 1.67 | k= 1, v=60 |
| 0.19 | 2.438 | 1.188 | | 1.75 | k= 2, v=60 |
| 0.56 | 2.061 | 2.918 | * | 1.77 | k= 3, v=60 |
| 1.67 | 1.443 | 5.759 | * | 1.78 | k= 4, v=60 |
| 5 | 1.084 | 7.412 | * | 1.79 | k= 5, v=60 |
| 15 | 1.023 | 6.980 | * | 1.79 | k= 6, v=60 |

s = 0.486

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 0.025 | 0.0029 | 0.22 | 0.47 | 0.12 |
| EC10 | 0.074 | 0.012 | 0.45 | 0.39 | 0.17 |
| EC25 | 0.46 | 0.14 | 1.5 | 0.26 | 0.30 |
| EC50 | 3.5 | 1.8 | 7.0 | 0.15 | 0.51 |

Slope = 0.765 Std.Err. = 0.137

!!!Poor fit: p = 0.016 based on DF= 4.0 60.

5701CW : cucumber dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|------|--------|--------------|---------------|----------------|-------------------|---------|
| | | | | | | |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | | |
|--------|------|------|-------|---------|------|------|
| 0.00 | 10.0 | 2.70 | 2.71 | -0.0167 | 100. | 0.00 |
| 0.0600 | 10.0 | 2.23 | 2.47 | -0.245 | 91.2 | 8.80 |
| 0.190 | 10.0 | 2.65 | 2.26 | 0.383 | 83.4 | 16.6 |
| 0.560 | 10.0 | 2.06 | 1.98 | 0.0825 | 72.9 | 27.1 |
| 1.67 | 10.0 | 1.44 | 1.62 | -0.179 | 59.8 | 40.2 |
| 5.00 | 10.0 | 1.08 | 1.23 | -0.147 | 45.4 | 54.6 |
| 15.0 | 7.00 | 1.02 | 0.855 | 0.168 | 31.5 | 68.5 |

!!!Warning: EC5 not bracketed by doses evaluated.

lettuce length

File: 570111 Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|----------|---------|--------|
| Between | 6 | 1620.570 | 270.095 | 95.037 |
| Within (Error) | 63 | 179.025 | 2.842 | |
| Total | 69 | 1799.595 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

lettuce length

File: 570111 Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|----------|-----|
| 1 | control | 16.680 | 16.680 | | |
| 2 | 0.06 | 15.780 | 15.780 | 1.194 | |
| 3 | 0.19 | 16.520 | 16.520 | 0.212 | |
| 4 | 0.56 | 13.790 | 13.790 | 3.833 * | |
| 5 | 1.67 | 7.940 | 7.940 | 11.593 * | |
| 6 | 5 | 5.500 | 5.500 | 14.829 * | |
| 7 | 15 | 5.340 | 5.340 | 15.041 * | |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

lettuce length

File: 570111 Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 2 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 1.772 | 10.6 | 0.900 |
| 3 | 0.19 | 10 | 1.772 | 10.6 | 0.160 |
| 4 | 0.56 | 10 | 1.772 | 10.6 | 2.890 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | |
|---|------|----|-------|------|--------|
| 5 | 1.67 | 10 | 1.772 | 10.6 | 8.740 |
| 6 | 5 | 10 | 1.772 | 10.6 | 11.180 |
| 7 | 15 | 10 | 1.772 | 10.6 | 11.340 |

lettuce length

File: 570111 Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 16.680 | 16.680 | 16.680 |
| 2 | 0.06 | 10 | 15.780 | 15.780 | 16.150 |
| 3 | 0.19 | 10 | 16.520 | 16.520 | 16.150 |
| 4 | 0.56 | 10 | 13.790 | 13.790 | 13.790 |
| 5 | 1.67 | 10 | 7.940 | 7.940 | 7.940 |
| 6 | 5 | 10 | 5.500 | 5.500 | 5.500 |
| 7 | 15 | 10 | 5.340 | 5.340 | 5.340 |

lettuce length

File: 570111 Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 16.680 | | | 1.67 | k= 1, v=63 |
| 0.06 | 16.150 | 0.703 | | 1.75 | k= 2, v=63 |
| 0.19 | 16.150 | 0.703 | * | 1.77 | k= 3, v=63 |
| 0.56 | 13.790 | 3.834 | * | 1.78 | k= 4, v=63 |
| 1.67 | 7.940 | 11.593 | * | 1.79 | k= 5, v=63 |
| 5 | 5.500 | 14.830 | * | 1.79 | k= 6, v=63 |
| 15 | 5.340 | 15.042 | * | 1.79 | |

s = 1.686

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.033 | 0.0094 | 0.12 | 0.27 | 0.29 |
| EC10 | 0.085 | 0.029 | 0.24 | 0.23 | 0.35 |
| EC25 | 0.41 | 0.20 | 0.87 | 0.16 | 0.48 |
| EC50 | 2.4 | 1.5 | 3.7 | 0.096 | 0.64 |

Slope = 0.883 Std.Err. = 0.0957

!!!Poor fit: p < 0.001 based on DF= 4.00 63.0

5701LL : lettuce length

Observed vs. Predicted Treatment Group Means

DP Barcode: D295614

MRID No.: 45745701

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 16.7 | 17.5 | -0.845 | 100. | 0.00 |
| 0.0600 | 10.0 | 15.8 | 16.1 | -0.367 | 92.1 | 7.87 |
| 0.190 | 10.0 | 16.5 | 14.6 | 1.89 | 83.5 | 16.5 |
| 0.560 | 10.0 | 13.8 | 12.5 | 1.32 | 71.2 | 28.8 |
| 1.67 | 10.0 | 7.94 | 9.73 | -1.79 | 55.5 | 44.5 |
| 5.00 | 10.0 | 5.50 | 6.82 | -1.32 | 38.9 | 61.1 |
| 15.0 | 10.0 | 5.34 | 4.23 | 1.11 | 24.1 | 75.9 |

!!!Warning: EC5 not bracketed by doses evaluated.

lettuce dry weight

File: 57011w Transform: SQUARE ROOT(Y)

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|-------|-------|--------|
| Between | 6 | 5.218 | 0.870 | 48.333 |
| Within (Error) | 63 | 1.161 | 0.018 | |
| Total | 69 | 6.379 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

lettuce dry weight

File: 57011w Transform: SQUARE ROOT(Y)

DUNNETTS TEST - TABLE 1 OF 2 Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|--------|-----|
| 1 | control | 1.100 | 1.234 | | |
| 2 | 0.06 | 1.029 | 1.088 | 1.190 | |
| 3 | 0.19 | 1.066 | 1.161 | 0.573 | |
| 4 | 0.56 | 0.886 | 0.799 | 3.572 | * |
| 5 | 1.67 | 0.603 | 0.367 | 8.292 | * |
| 6 | 5 | 0.482 | 0.242 | 10.300 | * |
| 7 | 15 | 0.398 | 0.170 | 11.703 | * |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

lettuce dry weight

File: 57011w Transform: SQUARE ROOT(Y)

DUNNETTS TEST - TABLE 2 OF 2 Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|----------------|--------------------------------------|-----------------|----------------------------|
| | | | | | |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | |
|---|---------|----|-------|------|-------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.290 | 23.5 | 0.146 |
| 3 | 0.19 | 10 | 0.290 | 23.5 | 0.073 |
| 4 | 0.56 | 10 | 0.290 | 23.5 | 0.435 |
| 5 | 1.67 | 10 | 0.290 | 23.5 | 0.867 |
| 6 | 5 | 10 | 0.290 | 23.5 | 0.992 |
| 7 | 15 | 10 | 0.290 | 23.5 | 1.064 |

lettuce dry weight

File: 5701lw Transform: SQUARE ROOT(Y)

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 1.234 | 1.100 | 1.100 |
| 2 | 0.06 | 10 | 1.088 | 1.029 | 1.047 |
| 3 | 0.19 | 10 | 1.161 | 1.066 | 1.047 |
| 4 | 0.56 | 10 | 0.799 | 0.886 | 0.886 |
| 5 | 1.67 | 10 | 0.367 | 0.603 | 0.603 |
| 6 | 5 | 10 | 0.242 | 0.482 | 0.482 |
| 7 | 15 | 10 | 0.170 | 0.398 | 0.398 |

lettuce dry weight

File: 5701lw Transform: SQUARE ROOT(Y)

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 1.100 | | | | |
| 0.06 | 1.047 | 0.872 | | 1.67 | k= 1, v=63 |
| 0.19 | 1.047 | 0.872 | | 1.75 | k= 2, v=63 |
| 0.56 | 0.886 | 3.531 | * | 1.77 | k= 3, v=63 |
| 1.67 | 0.603 | 8.197 | * | 1.78 | k= 4, v=63 |
| 5 | 0.482 | 10.182 | * | 1.79 | k= 5, v=63 |
| 15 | 0.398 | 11.569 | * | 1.79 | k= 6, v=63 |

s = 0.136

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.028 | 0.0067 | 0.11 | 0.31 | 0.24 |
| EC10 | 0.060 | 0.018 | 0.20 | 0.27 | 0.29 |
| EC25 | 0.22 | 0.087 | 0.55 | 0.20 | 0.40 |
| EC50 | 0.92 | 0.50 | 1.7 | 0.13 | 0.54 |

Slope = 1.08 Std.Err. = 0.139

!!!Poor fit: p = 0.025 based on DF= 4.0 63.

5701LW : lettuce dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 1.23 | 1.28 | -0.0437 | 100. | 0.00 |
| 0.0600 | 10.0 | 1.09 | 1.15 | -0.0620 | 90.0 | 10.0 |
| 0.190 | 10.0 | 1.16 | 0.984 | 0.177 | 77.0 | 23.0 |
| 0.560 | 10.0 | 0.799 | 0.757 | 0.0427 | 59.2 | 40.8 |
| 1.67 | 10.0 | 0.367 | 0.498 | -0.131 | 39.0 | 61.0 |
| 5.00 | 10.0 | 0.242 | 0.273 | -0.0311 | 21.4 | 78.6 |
| 15.0 | 10.0 | 0.170 | 0.122 | 0.0482 | 9.52 | 90.5 |

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC10 not bracketed by doses evaluated.

onion length

File: 5701ol Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|----------|---------|--------|
| Between | 6 | 1291.474 | 215.246 | 16.665 |
| Within (Error) | 61 | 787.871 | 12.916 | |
| Total | 67 | 2079.345 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

onion length

File: 5701ol Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 1 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|---------|-----|
| 1 | control | 25.060 | 25.060 | | |
| 2 | 0.06 | 25.000 | 25.000 | 0.037 | |
| 3 | 0.19 | 23.330 | 23.330 | 1.076 | |
| 4 | 0.56 | 24.410 | 24.410 | 0.404 | |
| 5 | 1.67 | 22.520 | 22.520 | 1.580 | |
| 6 | 5 | 15.370 | 15.370 | 6.029 * | |
| 7 | 15 | 13.300 | 13.300 | 6.898 * | |

Bonferroni T table value = 2.46 (1 Tailed Value, P=0.05, df=60, 6)

onion length

File: 5701ol Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 2 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 3.959 | 15.8 | 0.060 |
| 3 | 0.19 | 10 | 3.959 | 15.8 | 1.730 |
| 4 | 0.56 | 10 | 3.959 | 15.8 | 0.650 |
| 5 | 1.67 | 10 | 3.959 | 15.8 | 2.540 |
| 6 | 5 | 10 | 3.959 | 15.8 | 9.690 |
| 7 | 15 | 8 | 4.199 | 16.8 | 11.760 |

onion length

File: 5701ol

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 25.060 | 25.060 | 25.060 |
| 2 | 0.06 | 10 | 25.000 | 25.000 | 25.000 |
| 3 | 0.19 | 10 | 23.330 | 23.330 | 23.870 |
| 4 | 0.56 | 10 | 24.410 | 24.410 | 23.870 |
| 5 | 1.67 | 10 | 22.520 | 22.520 | 22.520 |
| 6 | 5 | 10 | 15.370 | 15.370 | 15.370 |
| 7 | 15 | 8 | 13.300 | 13.300 | 13.300 |

onion length

File: 5701ol

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 25.060 | | | | |
| 0.06 | 25.000 | 0.037 | | 1.67 | k= 1, v=61 |
| 0.19 | 23.870 | 0.740 | | 1.75 | k= 2, v=61 |
| 0.56 | 23.870 | 0.740 | | 1.77 | k= 3, v=61 |
| 1.67 | 22.520 | 1.580 | | 1.78 | k= 4, v=61 |
| 5 | 15.370 | 6.029 | * | 1.79 | k= 5, v=61 |
| 15 | 13.300 | 6.898 | * | 1.79 | k= 6, v=61 |

s = 3.594

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.40 | 0.10 | 1.6 | 0.29 | 0.26 |
| EC10 | 0.89 | 0.31 | 2.6 | 0.23 | 0.35 |
| EC25 | 3.4 | 1.9 | 6.2 | 0.13 | 0.55 |

DP Barcode: D295614

MRID No.: 45745701

EC50 15. 9.9 22. 0.089 0.67

Slope = 1.05 Std.Err. = 0.208

Goodness of fit: p = 0.11 based on DF= 4.0 61.

5701OL : onion length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 25.1 | 25.1 | -0.0259 | 100. | 0.00 |
| 0.0600 | 10.0 | 25.0 | 24.9 | 0.0644 | 99.4 | 0.599 |
| 0.190 | 10.0 | 23.3 | 24.5 | -1.17 | 97.7 | 2.34 |
| 0.560 | 10.0 | 24.4 | 23.4 | 1.02 | 93.3 | 6.75 |
| 1.67 | 10.0 | 22.5 | 21.1 | 1.43 | 84.1 | 15.9 |
| 5.00 | 10.0 | 15.4 | 17.3 | -1.95 | 69.1 | 30.9 |
| 15.0 | 8.00 | 13.3 | 12.5 | 0.790 | 49.9 | 50.1 |

onion dry weight

File: 5701ow Transform: 1/(SQUARE ROOT(Y))

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|--------|-------|--------|
| Between | 6 | 40.725 | 6.788 | 18.496 |
| Within (Error) | 61 | 22.374 | 0.367 | |
| Total | 67 | 63.099 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

onion dry weight

File: 5701ow Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST - TABLE 1 OF 2 Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|--------|-----|
| 1 | control | 2.851 | 0.141 | | |
| 2 | 0.06 | 2.864 | 0.139 | -0.051 | |
| 3 | 0.19 | 2.781 | 0.135 | 0.257 | |
| 4 | 0.56 | 2.718 | 0.152 | 0.488 | |
| 5 | 1.67 | 3.064 | 0.120 | -0.788 | |
| 6 | 5 | 4.011 | 0.065 | -4.284 | |
| 7 | 15 | 5.072 | 0.042 | -7.731 | |

Bonferroni T table value = 2.46 (1 Tailed Value, P=0.05, df=60, 6)

DP Barcode: D295614

MRID No.: 45745701

onion dry weight

File: 5701ow

Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST - TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | -0.087 | -61.3 | 0.002 |
| 3 | 0.19 | 10 | -0.087 | -61.3 | 0.006 |
| 4 | 0.56 | 10 | -0.087 | -61.3 | -0.010 |
| 5 | 1.67 | 10 | -0.087 | -61.3 | 0.022 |
| 6 | | 5 | -0.087 | -61.3 | 0.076 |
| 7 | | 15 | -0.095 | -67.0 | 0.099 |

onion dry weight

File: 5701ow

Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 0.141 | 2.851 | 2.804 |
| 2 | 0.06 | 10 | 0.139 | 2.864 | 2.804 |
| 3 | 0.19 | 10 | 0.135 | 2.781 | 2.804 |
| 4 | 0.56 | 10 | 0.152 | 2.718 | 2.804 |
| 5 | 1.67 | 10 | 0.120 | 3.064 | 3.064 |
| 6 | | 5 | 0.065 | 4.011 | 4.011 |
| 7 | | 15 | 0.042 | 5.072 | 5.072 |

onion dry weight

File: 5701ow

Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 2.804 | | | | |
| 0.06 | 2.804 | 0.174 | | 1.67 | k= 1, v=61 |
| 0.19 | 2.804 | 0.174 | | 1.75 | k= 2, v=61 |
| 0.56 | 2.804 | 0.174 | | 1.77 | k= 3, v=61 |
| 1.67 | 3.064 | 0.788 | | 1.78 | x= 4, v=61 |
| 5 | 4.011 | 4.286 | * | 1.79 | k= 5, v=61 |
| 15 | 5.072 | 7.733 | * | 1.79 | k= 6, v=61 |

s = 0.606

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | Std.Err. | Lower Bound / Estimate |
|-----------|----------|------------|----------|------------------------|
| | | Lower | Upper | |
| EC5 | 0.50 | 0.10 | 2.5 | 0.35 / 0.20 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | |
|------|------|------|-----|------|------|
| EC10 | 0.86 | 0.23 | 3.3 | 0.29 | 0.26 |
| EC25 | 2.1 | 0.86 | 5.2 | 0.20 | 0.41 |
| EC50 | 5.7 | 3.4 | 9.7 | 0.11 | 0.59 |

Slope = 1.56 Std.Err. = 0.422

Goodness of fit: p = 0.64 based on DF= 4.0 61.

5701OW : onion dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 0.141 | 0.144 | -0.00232 | 100. | 0.00 |
| 0.0600 | 10.0 | 0.139 | 0.144 | -0.00421 | 99.9 | 0.101 |
| 0.190 | 10.0 | 0.135 | 0.142 | -0.00698 | 98.9 | 1.05 |
| 0.560 | 10.0 | 0.152 | 0.135 | 0.0162 | 94.2 | 5.77 |
| 1.67 | 10.0 | 0.120 | 0.115 | 0.00495 | 79.8 | 20.2 |
| 5.00 | 10.0 | 0.0655 | 0.0772 | -0.0117 | 53.7 | 46.3 |
| 15.0 | 8.00 | 0.0421 | 0.0370 | 0.00510 | 25.8 | 74.2 |

oat length

File: 5701al Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|-----------|----------|---------|
| Between | 6 | 11493.288 | 1915.548 | 100.543 |
| Within (Error) | 59 | 1124.097 | 19.052 | |
| Total | 65 | 12617.385 | | |

Critical F value = 2.34 (0.05, 6, 40)

Since F > Critical F REJECT Ho:All groups equal

oat length

File: 5701al Transform: NO TRANSFORMATION

BONFERRONI T-TEST

TABLE 1 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|----------|-----|
| 1 | control | 48.540 | 48.540 | | |
| 2 | 0.06 | 45.660 | 45.660 | 1.475 | |
| 3 | 0.19 | 49.370 | 49.370 | -0.425 | |
| 4 | 0.56 | 47.900 | 47.900 | 0.328 | |
| 5 | 1.67 | 30.580 | 30.580 | 9.201 * | |
| 6 | 5 | 18.660 | 18.660 | 15.307 * | |
| 7 | 15 | 14.783 | 14.783 | 14.976 * | |

Bonferroni T table value = 2.48 (1 Tailed Value, P=0.05, df=50, 6)

oat length
File: 5701al Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 2 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 4.837 | 10.0 | 2.880 |
| 3 | 0.19 | 10 | 4.837 | 10.0 | -0.830 |
| 4 | 0.56 | 10 | 4.837 | 10.0 | 0.640 |
| 5 | 1.67 | 10 | 4.837 | 10.0 | 17.960 |
| 6 | 5 | 10 | 4.837 | 10.0 | 29.880 |
| 7 | 15 | 6 | 5.585 | 11.5 | 33.757 |

oat length
File: 5701al Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 48.540 | 48.540 | 48.540 |
| 2 | 0.06 | 10 | 45.660 | 45.660 | 47.643 |
| 3 | 0.19 | 10 | 49.370 | 49.370 | 47.643 |
| 4 | 0.56 | 10 | 47.900 | 47.900 | 47.643 |
| 5 | 1.67 | 10 | 30.580 | 30.580 | 30.580 |
| 6 | 5 | 10 | 18.660 | 18.660 | 18.660 |
| 7 | 15 | 6 | 14.783 | 14.783 | 14.783 |

oat length
File: 5701al Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 48.540 | | | | |
| 0.06 | 47.643 | 0.459 | | 1.68 | k= 1, v=59 |
| 0.19 | 47.643 | 0.459 | | 1.76 | k= 2, v=59 |
| 0.56 | 47.643 | 0.459 | | 1.79 | k= 3, v=59 |
| 1.67 | 30.580 | 9.201 | * | 1.80 | k= 4, v=59 |
| 5 | 18.660 | 15.307 | * | 1.80 | k= 5, v=59 |
| 15 | 14.783 | 14.976 | * | 1.81 | k= 6, v=59 |

s = 4.365

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | Std.Err. | Lower Bound |
|-----------|----------|------------|----------|-------------|
|-----------|----------|------------|----------|-------------|

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DP Barcode: D295614

MRID No.: 45745701

| | Lower | Upper | | /Estimate |
|------|-------|-------|------|-----------|
| EC5 | 0.16 | 0.078 | 0.35 | 0.16 |
| EC10 | 0.33 | 0.18 | 0.62 | 0.14 |
| EC25 | 1.0 | 0.68 | 1.6 | 0.093 |
| EC50 | 3.8 | 3.0 | 4.8 | 0.054 |

Slope = 1.21 Std.Err. = 0.116

!!!Poor fit: p < 0.001 based on DF= 4.00 59.0

5701AL : oat length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change %Control |
|--------|--------|--------------|---------------|----------------|-------------------|---------------------|
| 0.00 | 10.0 | 48.5 | 49.7 | -1.20 | 100. | 0.00 |
| 0.0600 | 10.0 | 45.7 | 49.0 | -3.35 | 98.5 | 1.48 |
| 0.190 | 10.0 | 49.4 | 46.8 | 2.52 | 94.2 | 5.82 |
| 0.560 | 10.0 | 47.9 | 41.9 | 6.02 | 84.2 | 15.8 |
| 1.67 | 10.0 | 30.6 | 33.1 | -2.55 | 66.6 | 33.4 |
| 5.00 | 10.0 | 18.7 | 22.0 | -3.31 | 44.2 | 55.8 |
| 15.0 | 6.00 | 14.8 | 11.7 | 3.11 | 23.5 | 76.5 |

oat dry weight

File: 5701aw Transform: NATURAL LOG(Y)

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|--------|-------|---------|
| Between | 6 | 41.847 | 6.975 | 139.500 |
| Within (Error) | 59 | 2.945 | 0.050 | |
| Total | 65 | 44.792 | | |

Critical F value = 2.34 (0.05, 6, 40)

Since F > Critical F REJECT Ho: All groups equal

oat dry weight

File: 5701aw Transform: NATURAL LOG(Y)

BONFERRONI T-TEST

TABLE 1 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|----------|-----|
| 1 | control | 0.266 | 1.320 | | |
| 2 | 0.06 | 0.235 | 1.281 | 0.312 | |
| 3 | 0.19 | 0.336 | 1.433 | -0.692 | |
| 4 | 0.56 | 0.256 | 1.324 | 0.103 | |
| 5 | 1.67 | -0.653 | 0.547 | 9.190 * | |
| 6 | 5 | -1.455 | 0.235 | 17.213 * | |
| 7 | 15 | -1.832 | 0.163 | 18.175 * | |

DP Barcode: D295614

MRID No.: 45745701

Bonferroni T table value = 2.48 (1 Tailed Value, P=0.05, df=50, 6)

oat dry weight

File: 5701aw Transform: NATURAL LOG(Y)

BONFERRONI T-TEST - TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.287 | 21.7 | 0.039 |
| 3 | 0.19 | 10 | 0.287 | 21.7 | -0.113 |
| 4 | 0.56 | 10 | 0.287 | 21.7 | -0.005 |
| 5 | 1.67 | 10 | 0.287 | 21.7 | 0.773 |
| 6 | 5 | 10 | 0.287 | 21.7 | 1.085 |
| 7 | 15 | 6 | 0.325 | 24.6 | 1.157 |

oat dry weight

File: 5701aw Transform: NATURAL LOG(Y)

WILLIAMS TEST (Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 1.320 | 0.266 | 0.279 |
| 2 | 0.06 | 10 | 1.281 | 0.235 | 0.279 |
| 3 | 0.19 | 10 | 1.433 | 0.336 | 0.279 |
| 4 | 0.56 | 10 | 1.324 | 0.256 | 0.256 |
| 5 | 1.67 | 10 | 0.547 | -0.653 | -0.653 |
| 6 | 5 | 10 | 0.235 | -1.455 | -1.455 |
| 7 | 15 | 6 | 0.163 | -1.832 | -1.832 |

oat dry weight

File: 5701aw Transform: NATURAL LOG(Y)

WILLIAMS TEST (Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 0.279 | | | | |
| 0.06 | 0.279 | 0.127 | | 1.68 | k= 1, v=59 |
| 0.19 | 0.279 | 0.127 | | 1.76 | k= 2, v=59 |
| 0.56 | 0.256 | 0.103 | | 1.79 | k= 3, v=59 |
| 1.67 | -0.653 | 9.197 | * | 1.80 | k= 4, v=59 |
| 5 | -1.455 | 17.227 | * | 1.80 | k= 5, v=59 |
| 15 | -1.832 | 18.190 | * | 1.81 | k= 6, v=59 |

s = 0.223

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.16 | 0.074 | 0.35 | 0.17 | 0.46 |
| EC10 | 0.27 | 0.14 | 0.52 | 0.14 | 0.51 |
| EC25 | 0.63 | 0.39 | 1.0 | 0.11 | 0.61 |
| EC50 | 1.6 | 1.2 | 2.3 | 0.070 | 0.72 |

Slope = 1.63 Std.Err. = 0.182

!!!Poor fit: p < 0.001 based on DF= 4.00 59.0

5701AW : oat dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 1.32 | 1.40 | -0.0830 | 100. | 0.00 |
| 0.0600 | 10.0 | 1.28 | 1.39 | -0.109 | 99.0 | 0.962 |
| 0.190 | 10.0 | 1.43 | 1.31 | 0.120 | 93.6 | 6.38 |
| 0.560 | 10.0 | 1.32 | 1.09 | 0.236 | 77.6 | 22.4 |
| 1.67 | 10.0 | 0.547 | 0.692 | -0.145 | 49.3 | 50.7 |
| 5.00 | 10.0 | 0.235 | 0.300 | -0.0648 | 21.4 | 78.6 |
| 15.0 | 6.00 | 0.163 | 0.0813 | 0.0822 | 5.79 | 94.2 |

radish length

File: 5701rl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|---------|---------|--------|
| Between | 5 | 722.031 | 144.406 | 47.896 |
| Within (Error) | 50 | 150.725 | 3.015 | |
| Total | 55 | 872.756 | | |

Critical F value = 2.45 (0.05,5,40)

Since F > Critical F REJECT Ho:All groups equal

radish length

File: 5701rl Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 1 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|--------|-----|
| 1 | control | 13.970 | 13.970 | | |
| 2 | 0.06 | 14.700 | 14.700 | -0.940 | |
| 3 | 0.19 | 15.300 | 15.300 | -1.713 | |
| 4 | 0.56 | 9.900 | 9.900 | 5.241 | * |
| 5 | 1.67 | 7.260 | 7.260 | 8.641 | * |
| 6 | 5 | 5.600 | 5.600 | 9.335 | * |

DP Barcode: D295614

MRID No.: 45745701

Bonferroni T table value = 2.40 (1 Tailed Value, P=0.05, df=50,5)

radish length
 File: 5701rl Transform: NO TRANSFORMATION

BONFERRONI T-TEST

TABLE 2 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 1.867 | 13.4 | -0.730 |
| 3 | 0.19 | 10 | 1.867 | 13.4 | -1.330 |
| 4 | 0.56 | 10 | 1.867 | 13.4 | 4.070 |
| 5 | 1.67 | 10 | 1.867 | 13.4 | 6.710 |
| 6 | | 5 6 | 2.156 | 15.4 | 8.370 |

radish length
 File: 5701rl Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|-----|---------------|------------------|-----------------|
| 1 | control | 10 | 13.970 | 13.970 | 14.657 |
| 2 | 0.06 | 10 | 14.700 | 14.700 | 14.657 |
| 3 | 0.19 | 10 | 15.300 | 15.300 | 14.657 |
| 4 | 0.56 | 10 | 9.900 | 9.900 | 9.900 |
| 5 | 1.67 | 10 | 7.260 | 7.260 | 7.260 |
| 6 | | 5 6 | 5.600 | 5.600 | 5.600 |

radish length
 File: 5701rl Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 14.657 | | | | |
| 0.06 | 14.657 | 0.884 | | 1.68 | k= 1, v=50 |
| 0.19 | 14.657 | 0.884 | | 1.76 | k= 2, v=50 |
| 0.56 | 9.900 | 5.242 | * | 1.79 | k= 3, v=50 |
| 1.67 | 7.260 | 8.642 | * | 1.80 | k= 4, v=50 |
| 5 | 5.600 | 9.335 | * | 1.80 | k= 5, v=50 |

S = 1.736

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | Std.Err. | Lower Bound |
|-----------|----------|------------|----------|-------------|
|-----------|----------|------------|----------|-------------|

DP Barcode: D295614

MRID No.: 45745701

| | | Lower | Upper | /Estimate |
|------|-------|-------|-------|-----------|
| EC5 | 0.074 | 0.023 | 0.24 | 0.26 |
| EC10 | 0.15 | 0.058 | 0.40 | 0.21 |
| EC25 | 0.51 | 0.27 | 0.97 | 0.14 |
| EC50 | 2.0 | 1.4 | 2.8 | 0.078 |

Slope = 1.16 Std.Err. = 0.177

!!!Poor fit: p < 0.001 based on DF= 3.00 50.0

5701RL : radish length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 14.0 | 15.0 | -1.03 | 100. | 0.00 |
| 0.0600 | 10.0 | 14.7 | 14.4 | 0.301 | 96.0 | 4.00 |
| 0.190 | 10.0 | 15.3 | 13.2 | 2.11 | 87.9 | 12.1 |
| 0.560 | 10.0 | 9.90 | 11.0 | -1.13 | 73.6 | 26.4 |
| 1.67 | 10.0 | 7.26 | 7.99 | -0.729 | 53.3 | 46.7 |
| 5.00 | 6.00 | 5.60 | 4.80 | 0.803 | 32.0 | 68.0 |

radish dry weight

File: 5701rw Transform: 1/(SQUARE ROOT(Y))

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|-------|-------|--------|
| Between | 5 | 3.395 | 0.679 | 33.950 |
| Within (Error) | 50 | 1.022 | 0.020 | |
| Total | 55 | 4.417 | | |

Critical F value = 2.45 (0.05,5,40)

Since F > Critical F REJECT Ho:All groups equal

radish dry weight

File: 5701rw Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST - TABLE 1 OF 2 Ho:Control<Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|--------|-----|
| 1 | control | 1.057 | 0.958 | | |
| 2 | 0.06 | 1.000 | 1.009 | 0.890 | |
| 3 | 0.19 | 1.012 | 1.011 | 0.703 | |
| 4 | 0.56 | 1.367 | 0.545 | -4.914 | |
| 5 | 1.67 | 1.583 | 0.409 | -8.321 | |
| 6 | 5 | 1.563 | 0.429 | -6.939 | |

Bonferroni T table value = 2.40 (1 Tailed Value, P=0.05, df=50,5)

radish dry weight
File: 5701rw

Transform: 1/(SQUARE ROOT(Y))

BONFERRONI T-TEST - TABLE 2 OF 2 Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN. ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|------------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | -0.326 | -34.1 | -0.050 |
| 3 | 0.19 | 10 | -0.326 | -34.1 | -0.053 |
| 4 | 0.56 | 10 | -0.326 | -34.1 | 0.414 |
| 5 | 1.67 | 10 | -0.326 | -34.1 | 0.550 |
| 6 | 5 | 6 | -0.393 | -41.0 | 0.529 |

radish dry weight

File: 5701rw

Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 0.958 | 1.057 | 1.023 |
| 2 | 0.06 | 10 | 1.009 | 1.000 | 1.023 |
| 3 | 0.19 | 10 | 1.011 | 1.012 | 1.023 |
| 4 | 0.56 | 10 | 0.545 | 1.367 | 1.367 |
| 5 | 1.67 | 10 | 0.409 | 1.583 | 1.576 |
| 6 | 5 | 6 | 0.429 | 1.563 | 1.576 |

radish dry weight

File: 5701rw

Transform: 1/(SQUARE ROOT(Y))

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG WILLIAMS | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|--------------|----------------|--------------------|
| control | 1.023 | | | | |
| 0.06 | 1.023 | 0.526 | | 1.68 | k= 1, v=50 |
| 0.19 | 1.023 | 0.526 | | 1.76 | k= 2, v=50 |
| 0.56 | 1.367 | 4.862 | * | 1.79 | k= 3, v=50 |
| 1.67 | 1.576 | 8.119 | * | 1.80 | k= 4, v=50 |
| 5 | 1.576 | 7.031 | * | 1.80 | k= 5, v=50 |

s = 0.143

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|----------|-----------------------|
| | | Lower | Upper | |
| EC5 | 0.028 | 0.0045 | 0.18 | 0.40 0.16 |
| EC10 | 0.068 | 0.015 | 0.31 | 0.33 0.22 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | |
|------|------|------|------|------|------|
| EC25 | 0.29 | 0.10 | 0.81 | 0.22 | 0.36 |
| EC50 | 1.5 | 0.81 | 2.6 | 0.13 | 0.55 |

$$\text{Slope} = 0.962 \quad \text{Std.Err.} = 0.187$$

!!!Poor fit: p < 0.001 based on DF= 3.00 50.0

5701RW : radish dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 0.958 | 1.04 | -0.0818 | 100. | 0.00 |
| 0.0600 | 10.0 | 1.01 | 0.945 | 0.0633 | 90.9 | 9.12 |
| 0.190 | 10.0 | 1.01 | 0.835 | 0.176 | 80.3 | 19.7 |
| 0.560 | 10.0 | 0.545 | 0.682 | -0.137 | 65.5 | 34.5 |
| 1.67 | 10.0 | 0.409 | 0.497 | -0.0881 | 47.7 | 52.3 |
| 5.00 | 6.00 | 0.429 | 0.316 | 0.113 | 30.3 | 69.7 |

!!!Warning: EC5 not bracketed by doses evaluated.

soybean length

File: 5701sl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|----------|---------|---------|
| Between | 6 | 4816.673 | 802.779 | 118.789 |
| Within (Error) | 63 | 425.745 | 6.758 | |
| Total | 69 | 5242.418 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho: All groups equal

soybean length

File: 5701sl Transform: NO TRANSFORMATION

DUNNETTS TEST

TABLE 1 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|--------|-----|
| 1 | control | 41.610 | 41.610 | | |
| 2 | 0.06 | 42.220 | 42.220 | -0.525 | |
| 3 | 0.19 | 41.520 | 41.520 | 0.077 | |
| 4 | 0.56 | 33.710 | 33.710 | 6.795 | * |
| 5 | 1.67 | 25.230 | 25.230 | 14.089 | * |
| 6 | 5 | 23.510 | 23.510 | 15.569 | * |
| 7 | 15 | 23.050 | 23.050 | 15.964 | * |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

soybean length
File: 5701sl

Transform: NO TRANSFORMATION

DUNNETTS TEST

TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 2.732 | 6.6 | -0.610 |
| 3 | 0.19 | 10 | 2.732 | 6.6 | 0.090 |
| 4 | 0.56 | 10 | 2.732 | 6.6 | 7.900 |
| 5 | 1.67 | 10 | 2.732 | 6.6 | 16.380 |
| 6 | 5 | 10 | 2.732 | 6.6 | 18.100 |
| 7 | 15 | 10 | 2.732 | 6.6 | 18.560 |

soybean length
File: 5701sl

Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 41.610 | 41.610 | 41.915 |
| 2 | 0.06 | 10 | 42.220 | 42.220 | 41.915 |
| 3 | 0.19 | 10 | 41.520 | 41.520 | 41.520 |
| 4 | 0.56 | 10 | 33.710 | 33.710 | 33.710 |
| 5 | 1.67 | 10 | 25.230 | 25.230 | 25.230 |
| 6 | 5 | 10 | 23.510 | 23.510 | 23.510 |
| 7 | 15 | 10 | 23.050 | 23.050 | 23.050 |

soybean length
File: 5701sl

Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 41.915 | | | | |
| 0.06 | 41.915 | 0.262 | | 1.67 | k= 1, v=63 |
| 0.19 | 41.520 | 0.077 | | 1.75 | k= 2, v=63 |
| 0.56 | 33.710 | 6.795 | * | 1.77 | k= 3, v=63 |
| 1.67 | 25.230 | 14.089 | * | 1.78 | k= 4, v=63 |
| 5 | 23.510 | 15.569 | * | 1.79 | k= 5, v=63 |
| 15 | 23.050 | 15.965 | * | 1.79 | k= 6, v=63 |

s = 2.600

Note: df used for table values are approximate when v > 20.

Estimates of ECT

| Parameter | Estimate | 95% Bounds | Std.Err. | Lower Bound |
|-----------|----------|------------|----------|-------------|
|-----------|----------|------------|----------|-------------|

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DP Barcode: D295614

MRID No.: 45745701

| | | Lower | Upper | /Estimate |
|------|-------|--------|-------|------------|
| EC5 | 0.019 | 0.0043 | 0.082 | 0.32 0.23 |
| EC10 | 0.076 | 0.023 | 0.25 | 0.26 0.31 |
| EC25 | 0.79 | 0.38 | 1.6 | 0.16 0.48 |
| EC50 | 11. | 7.0 | 16. | 0.092 0.66 |

Slope = 0.598 Std.Err. = 0.0655

!!!Poor fit: p < 0.001 based on DF= 4.00 63.0

5701SL : soybean length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 41.6 | 43.8 | -2.22 | 100. | 0.00 |
| 0.0600 | 10.0 | 42.2 | 39.9 | 2.31 | 91.1 | 8.95 |
| 0.190 | 10.0 | 41.5 | 37.3 | 4.18 | 85.2 | 14.8 |
| 0.560 | 10.0 | 33.7 | 34.1 | -0.376 | 77.8 | 22.2 |
| 1.67 | 10.0 | 25.2 | 30.0 | -4.78 | 68.5 | 31.5 |
| 5.00 | 10.0 | 23.5 | 25.3 | -1.81 | 57.8 | 42.2 |
| 15.0 | 10.0 | 23.1 | 20.4 | 2.69 | 46.5 | 53.5 |

!!!Warning: EC5 not bracketed by doses evaluated.

soybean dry weight

File: 5701sw Transform: SQUARE ROOT(Y)

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|--------|-------|---------|
| Between | 6 | 14.719 | 2.453 | 102.208 |
| Within (Error) | 63 | 1.497 | 0.024 | |
| Total | 69 | 16.216 | | |

Critical F value = 2.25 (0.05, 6, 60)

Since F > Critical F REJECT Ho:All groups equal

soybean dry weight

File: 5701sw Transform: SQUARE ROOT(Y)

DUNNETTS TEST - TABLE 1 OF 2

Ho:Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|---------------------|--------------------------------------|----------|-----|
| 1 | control | 2.068 | 4.299 | | |
| 2 | 0.06 | 2.052 | 4.256 | 0.238 | |
| 3 | 0.19 | 2.055 | 4.254 | 0.184 | |
| 4 | 0.56 | 1.764 | 3.131 | 4.385 * | |
| 5 | 1.67 | 1.229 | 1.525 | 12.111 * | |
| 6 | 5 | 0.993 | 1.000 | 15.522 * | |

AF

DP Barcode: D295614

MRID No.: 45745701

| | | | | |
|---|----|-------|-------|----------|
| 7 | 15 | 1.045 | 1.099 | 14.768 * |
|---|----|-------|-------|----------|

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

soybean dry weight

File: 5701sw Transform: SQUARE ROOT(Y)

DUNNETTS TEST

TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.647 | 15.1 | 0.043 |
| 3 | 0.19 | 10 | 0.647 | 15.1 | 0.045 |
| 4 | 0.56 | 10 | 0.647 | 15.1 | 1.168 |
| 5 | 1.67 | 10 | 0.647 | 15.1 | 2.773 |
| 6 | 5 | 10 | 0.647 | 15.1 | 3.298 |
| 7 | 15 | 10 | 0.647 | 15.1 | 3.200 |

soybean dry weight

File: 5701sw Transform: SQUARE ROOT(Y)

WILLIAMS TEST

(Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 4.299 | 2.068 | 2.068 |
| 2 | 0.06 | 10 | 4.256 | 2.052 | 2.054 |
| 3 | 0.19 | 10 | 4.254 | 2.055 | 2.054 |
| 4 | 0.56 | 10 | 3.131 | 1.764 | 1.764 |
| 5 | 1.67 | 10 | 1.525 | 1.229 | 1.229 |
| 6 | 5 | 10 | 1.000 | 0.993 | 1.019 |
| 7 | 15 | 10 | 1.099 | 1.045 | 1.019 |

soybean dry weight

File: 5701sw Transform: SQUARE ROOT(Y)

WILLIAMS TEST

(Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|-----------|----------------|--------------------|
| control | 2.068 | | | | |
| 0.06 | 2.054 | 0.212 | | 1.67 | k= 1, v=63 |
| 0.19 | 2.054 | 0.212 | | 1.75 | k= 2, v=63 |
| 0.56 | 1.764 | 4.407 | * | 1.77 | k= 3, v=63 |
| 1.67 | 1.229 | 12.169 | * | 1.78 | k= 4, v=63 |
| 5 | 1.019 | 15.218 | * | 1.79 | k= 5, v=63 |
| 15 | 1.019 | 15.218 | * | 1.79 | k= 6, v=63 |

s = 0.154

Note: df used for table values are approximate when v > 20.

DP Barcode: D295614

MRID No.: 45745701

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.021 | 0.0059 | 0.077 | 0.28 | 0.28 |
| EC10 | 0.052 | 0.017 | 0.16 | 0.24 | 0.33 |
| EC25 | 0.24 | 0.11 | 0.53 | 0.18 | 0.45 |
| EC50 | 1.3 | 0.76 | 2.1 | 0.11 | 0.60 |

Slope = 0.928 Std.Err. = 0.0995

!!!Poor fit: p < 0.001 based on DF= 4.00 63.0

5701SW : soybean dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|-----------|------------|-------------|----------------|---------|
| 0.00 | 10.0 | 4.30 | 4.66 | -0.359 | 100. | 0.00 |
| 0.0600 | 10.0 | 4.26 | 4.15 | 0.110 | 89.0 | 11.0 |
| 0.190 | 10.0 | 4.25 | 3.62 | 0.633 | 77.7 | 22.3 |
| 0.560 | 10.0 | 3.13 | 2.93 | 0.204 | 62.8 | 37.2 |
| 1.67 | 10.0 | 1.53 | 2.12 | -0.595 | 45.5 | 54.5 |
| 5.00 | 10.0 | 1.00 | 1.35 | -0.349 | 29.0 | 71.0 |
| 15.0 | 10.0 | 1.10 | 0.743 | 0.356 | 15.9 | 84.1 |

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC10 not bracketed by doses evaluated.

tomato length

File: 5701tl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|----------|----------|--------|
| Between | 6 | 6693.604 | 1115.601 | 95.155 |
| Within (Error) | 51 | 597.935 | 11.724 | |
| Total | 57 | 7291.539 | | |

Critical F value = 2.34 (0.05, 6, 40)

Since F > Critical F REJECT Ho: All groups equal

tomato length

File: 5701tl Transform: NO TRANSFORMATION

BONFERRONI T-TEST - TABLE 1 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | | |
|-------|----------------|------------------|-----------------------------------|-----|--|
| | | | T STAT | SIG | |
| 1 | control | 36.920 | 36.920 | | |

DP Barcode: D295614

MRID No.: 45745701

| | | | | |
|---|------|--------|--------|----------|
| 2 | 0.06 | 34.060 | 34.060 | 1.868 |
| 3 | 0.19 | 35.900 | 35.900 | 0.666 |
| 4 | 0.56 | 24.320 | 24.320 | 8.228 * |
| 5 | 1.67 | 14.560 | 14.560 | 14.602 * |
| 6 | 5 | 8.250 | 8.250 | 16.215 * |
| 7 | 15 | 8.000 | 8.000 | 10.904 * |

Bonferroni T table value = 2.48 (1 Tailed Value, P=0.05, df=50, 6)

tomato length

File: 5701tl

Transform: NO TRANSFORMATION

BONFERRONI T-TEST

TABLE 2 OF 2

Ho:Control<Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 3.794 | 10.3 | 2.860 |
| 3 | 0.19 | 10 | 3.794 | 10.3 | 1.020 |
| 4 | 0.56 | 10 | 3.794 | 10.3 | 12.600 |
| 5 | 1.67 | 10 | 3.794 | 10.3 | 22.360 |
| 6 | 5 | 6 | 4.382 | 11.9 | 28.670 |
| 7 | 15 | 2 | 6.572 | 17.8 | 28.920 |

tomato length

File: 5701tl

Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 36.920 | 36.920 | 36.920 |
| 2 | 0.06 | 10 | 34.060 | 34.060 | 34.980 |
| 3 | 0.19 | 10 | 35.900 | 35.900 | 34.980 |
| 4 | 0.56 | 10 | 24.320 | 24.320 | 24.320 |
| 5 | 1.67 | 10 | 14.560 | 14.560 | 14.560 |
| 6 | 5 | 6 | 8.250 | 8.250 | 8.250 |
| 7 | 15 | 2 | 8.000 | 8.000 | 8.000 |

tomato length

File: 5701tl

Transform: NO TRANSFORMATION

WILLIAMS TEST

(Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG. P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|-----------------|----------------|------------|----------------|--------------------|
| control | 36.920 | | | | |
| 0.06 | 34.980 | 1.267 | | 1.68 | k= 1, v=51 |
| 0.19 | 34.980 | 1.267 | | 1.76 | k= 2, v=51 |
| 0.56 | 24.320 | 8.228 | * | 1.79 | k= 3, v=51 |
| 1.67 | 14.560 | 14.602 | * | 1.80 | k= 4, v=51 |

DP Barcode: D295614

MRID No.: 45745701

| | | | | | |
|----|-------|--------|---|------|------------|
| 5 | 8.250 | 16.214 | * | 1.80 | k= 5, v=51 |
| 15 | 8.000 | 10.904 | * | 1.81 | k= 6, v=51 |

s = 3.424

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-------------|--------------|--------------|-------------|-------------|-----------------------|
| | | Lower | Upper | | |
| EC5 | 0.044 | 0.019 | 0.10 | 0.18 | 0.43 |
| EC10 | 0.091 | 0.045 | 0.19 | 0.15 | 0.49 |
| EC25 | 0.31 | 0.19 | 0.51 | 0.11 | 0.60 |
| EC50 | 1.2 | 0.88 | 1.6 | 0.068 | 0.73 |

Slope = 1.15 Std.Err. = 0.107

!!!Poor fit: p < 0.001 based on DF= 4.00 51.0

5701TL : tomato length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 36.9 | 38.1 | -1.16 | 100. | 0.00 |
| 0.0600 | 10.0 | 34.1 | 35.5 | -1.43 | 93.2 | 6.81 |
| 0.190 | 10.0 | 35.9 | 31.2 | 4.65 | 82.0 | 18.0 |
| 0.560 | 10.0 | 24.3 | 24.7 | -0.351 | 64.8 | 35.2 |
| 1.67 | 10.0 | 14.6 | 16.6 | -2.00 | 43.5 | 56.5 |
| 5.00 | 6.00 | 8.25 | 9.10 | -0.853 | 23.9 | 76.1 |
| 15.0 | 2.00 | 8.00 | 3.98 | 4.02 | 10.5 | 89.5 |

!!!Warning: EC5 not bracketed by doses evaluated.

tomato dry weight

File: 5701tw Transform: NO TRANSFORM

KRUSKAL-WALLIS ANOVA BY RANKS - TABLE 1 OF 2

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | RANK SUM |
|-------|----------------|---------------------|--------------------------------------|-------------|
| 1 | control | 2.843 | 2.843 | 445.000 |
| 2 | 0.06 | 2.484 | 2.484 | 409.000 |
| 3 | 0.19 | 2.711 | 2.711 | 436.000 |
| 4 | 0.56 | 1.287 | 1.287 | 247.000 |
| 5 | 1.67 | 0.402 | 0.402 | 127.000 |
| 6 | 5 | 0.218 | 0.218 | 31.000 |
| 7 | 15 | 0.285 | 0.285 | 16.000 |

Calculated H Value = 45.825 Critical H Value Table = 12.590
Since Calc H > Crit H REJECT Ho:All groups are equal.**tomato dry weight**

File: 5701tw Transform: NO TRANSFORM

DUNNS MULTIPLE COMPARISON - KRUSKAL-WALLIS - TABLE 2 OF 2

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | ORIGINAL MEAN | GROUP | | | | | |
|-------|----------------|---------------------|------------------|-------|---|---|---|---|---|
| | | | | 0 | 0 | 0 | 0 | 0 | 0 |
| 6 | 5 | 0.218 | 0.218 | \ | | | | | |
| 7 | 15 | 0.285 | 0.285 | \ | | | | | |
| 5 | 1.67 | 0.402 | 0.402 | | | | | | \ |
| 4 | 0.56 | 1.287 | 1.287 | | | | | | \ |
| 2 | 0.06 | 2.484 | 2.484 | * | * | * | * | * | \ |
| 3 | 0.19 | 2.711 | 2.711 | * | * | * | * | * | \ |
| 1 | control | 2.843 | 2.843 | * | * | * | * | * | \ |

* = significant difference ($p=0.05$)
 Table q value (0.05, 7) = 3.038

= no significant difference
 Unequal reps - multiple SE values

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 0.041 | 0.013 | 0.13 | 0.26 | 0.31 |
| EC10 | 0.071 | 0.025 | 0.20 | 0.22 | 0.36 |
| EC25 | 0.18 | 0.082 | 0.39 | 0.17 | 0.46 |
| EC50 | 0.50 | 0.29 | 0.83 | 0.11 | 0.59 |

Slope = 1.52 Std.Err. = 0.229

!!!Poor fit: $p < 0.001$ based on DF= 4.00 51.0

5701TW : tomato dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 2.84 | 2.94 | -0.0960 | 100. | 0.00 |
| 0.0600 | 10.0 | 2.48 | 2.70 | -0.214 | 91.8 | 8.19 |
| 0.190 | 10.0 | 2.71 | 2.16 | 0.548 | 73.6 | 26.4 |
| 0.560 | 10.0 | 1.29 | 1.37 | -0.0876 | 46.8 | 53.2 |
| 1.67 | 10.0 | 0.402 | 0.621 | -0.219 | 21.1 | 78.9 |
| 5.00 | 6.00 | 0.218 | 0.187 | 0.0317 | 6.35 | 93.6 |
| 15.0 | 2.00 | 0.285 | 0.0359 | 0.249 | 1.22 | 98.8 |

!!!Warning: EC5 not bracketed by doses evaluated.

wheat length

File: 5701wl Transform: NO TRANSFORMATION

ANOVA TABLE

| SOURCE | DF | SS | MS | F |
|----------------|----|---------|--------|--------|
| Between | 6 | 468.716 | 78.119 | 11.277 |
| Within (Error) | 63 | 436.382 | 6.927 | |

DP Barcode: D295614

MRID No.: 45745701

Total 69 905.098

Critical F value = 2.25 (0.05, 6, 60)
 Since F > Critical F REJECT Ho: All groups equal

wheat length
 File: 5701wl Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|---------|-----|
| 1 | control | 28.020 | 28.020 | | |
| 2 | 0.06 | 31.300 | 31.300 | -2.787 | |
| 3 | 0.19 | 29.720 | 29.720 | -1.444 | |
| 4 | 0.56 | 29.460 | 29.460 | -1.223 | |
| 5 | 1.67 | 28.160 | 28.160 | -0.119 | |
| 6 | 5 | 26.430 | 26.430 | 1.351 | |
| 7 | 15 | 22.670 | 22.670 | 4.545 * | |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

wheat length
 File: 5701wl Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 2 OF 2 Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 2.766 | 9.9 | -3.280 |
| 3 | 0.19 | 10 | 2.766 | 9.9 | -1.700 |
| 4 | 0.56 | 10 | 2.766 | 9.9 | -1.440 |
| 5 | 1.67 | 10 | 2.766 | 9.9 | -0.140 |
| 6 | 5 | 10 | 2.766 | 9.9 | 1.590 |
| 7 | 15 | 10 | 2.766 | 9.9 | 5.350 |

wheat length
 File: 5701wl Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 28.020 | 28.020 | 29.680 |
| 2 | 0.06 | 10 | 31.300 | 31.300 | 29.680 |
| 3 | 0.19 | 10 | 29.720 | 29.720 | 29.680 |
| 4 | 0.56 | 10 | 29.460 | 29.460 | 29.460 |
| 5 | 1.67 | 10 | 28.160 | 28.160 | 28.160 |
| 6 | 5 | 10 | 26.430 | 26.430 | 26.430 |
| 7 | 15 | 10 | 22.670 | 22.670 | 22.670 |

wheat length

File: 5701wl

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|--------------------|-------------------|--------------|-------------------|-----------------------|
| control | 29.680 | | | | |
| 0.06 | 29.680 | 1.410 | | 1.67 | k= 1, v=63 |
| 0.19 | 29.680 | 1.410 | | 1.75 | k= 2, v=63 |
| 0.56 | 29.460 | 1.223 | | 1.77 | k= 3, v=63 |
| 1.67 | 28.160 | 0.119 | | 1.78 | k= 4, v=63 |
| 5 | 26.430 | 1.351 | | 1.79 | k= 5, v=63 |
| 15 | 22.670 | 4.545 | * | 1.79 | k= 6, v=63 |

s = 2.632

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|---------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 1.9 | 0.63 | 6.0 | 0.25 | 0.32 |
| EC10 | 4.3 | 2.1 | 9.0 | 0.16 | 0.48 |
| EC25 | 16. | 11. | 24. | 0.086 | 0.67 |
| EC50 | 72. | 27. | 2.0E+02 | 0.22 | 0.37 |

Slope = 1.05 Std.Err. = 0.290

Goodness of fit: p = 0.12 based on DF= 4.0 63.

5701WL : wheat length

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|---------|
| 0.00 | 10.0 | 28.0 | 29.7 | -1.69 | 100. | 0.00 |
| 0.0600 | 10.0 | 31.3 | 29.7 | 1.61 | 99.9 | 0.0623 |
| 0.190 | 10.0 | 29.7 | 29.6 | 0.116 | 99.7 | 0.343 |
| 0.560 | 10.0 | 29.5 | 29.3 | 0.155 | 98.7 | 1.35 |
| 1.67 | 10.0 | 28.2 | 28.4 | -0.262 | 95.7 | 4.32 |
| 5.00 | 10.0 | 26.4 | 26.4 | 0.0540 | 88.8 | 11.2 |
| 15.0 | 10.0 | 22.7 | 22.7 | 0.00831 | 76.3 | 23.7 |

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

wheat dry weight

File: 5701ww

Transform: NO TRANSFORMATION

ANOVA TABLE

DP Barcode: D295614

MRID No.: 45745701

| SOURCE | DF | SS | MS | F |
|----------------|----|-------|-------|-------|
| Between | 6 | 1.346 | 0.224 | 5.895 |
| Within (Error) | 63 | 2.388 | 0.038 | |
| Total | 69 | 3.734 | | |

Critical F value = 2.25 (0.05, 6, 60)
 Since F > Critical F REJECT Ho: All groups equal

wheat dry weight

File: 5701ww Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | TRANSFORMED MEAN | MEAN CALCULATED IN ORIGINAL UNITS | T STAT | SIG |
|-------|----------------|------------------|-----------------------------------|---------|-----|
| 1 | control | 0.708 | 0.708 | | |
| 2 | 0.06 | 0.786 | 0.786 | -0.900 | |
| 3 | 0.19 | 0.873 | 0.873 | -1.893 | |
| 4 | 0.56 | 0.951 | 0.951 | -2.790 | |
| 5 | 1.67 | 0.791 | 0.791 | -0.950 | |
| 6 | 5 | 0.726 | 0.726 | -0.205 | |
| 7 | 15 | 0.477 | 0.477 | 2.643 * | |

Dunnett table value = 2.35 (1 Tailed Value, P=0.05, df=60, 6)

wheat dry weight

File: 5701ww Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 2 OF 2

Ho: Control < Treatment

| GROUP | IDENTIFICATION | NUM OF REPS | Minimum Sig Diff (IN ORIG. UNITS) | % of CONTROL | DIFFERENCE FROM CONTROL |
|-------|----------------|-------------|-----------------------------------|--------------|-------------------------|
| 1 | control | 10 | | | |
| 2 | 0.06 | 10 | 0.205 | 28.9 | -0.078 |
| 3 | 0.19 | 10 | 0.205 | 28.9 | -0.165 |
| 4 | 0.56 | 10 | 0.205 | 28.9 | -0.243 |
| 5 | 1.67 | 10 | 0.205 | 28.9 | -0.083 |
| 6 | 5 | 10 | 0.205 | 28.9 | -0.018 |
| 7 | 15 | 10 | 0.205 | 28.9 | 0.230 |

wheat dry weight

File: 5701ww Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model)

TABLE 1 OF 2

| GROUP | IDENTIFICATION | N | ORIGINAL MEAN | TRANSFORMED MEAN | ISOTONIZED MEAN |
|-------|----------------|----|---------------|------------------|-----------------|
| 1 | control | 10 | 0.708 | 0.708 | 0.829 |

36

| | | | | | |
|---|------|----|-------|-------|-------|
| 2 | 0.06 | 10 | 0.786 | 0.786 | 0.829 |
| 3 | 0.19 | 10 | 0.873 | 0.873 | 0.829 |
| 4 | 0.56 | 10 | 0.951 | 0.951 | 0.829 |
| 5 | 1.67 | 10 | 0.791 | 0.791 | 0.791 |
| 6 | 5 | 10 | 0.726 | 0.726 | 0.726 |
| 7 | 15 | 10 | 0.477 | 0.477 | 0.477 |

wheat dry weight

File: 5701ww

Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model)

TABLE 2 OF 2

| IDENTIFICATION | ISOTONIZED MEAN | CALC. WILLIAMS | SIG P=.05 | TABLE WILLIAMS | DEGREES OF FREEDOM |
|----------------|--------------------|-------------------|--------------|-------------------|-----------------------|
| control | 0.829 | | | | |
| 0.06 | 0.829 | 1.398 | | 1.67 | k= 1, v=63 |
| 0.19 | 0.829 | 1.398 | | 1.75 | k= 2, v=63 |
| 0.56 | 0.829 | 1.398 | | 1.77 | k= 3, v=63 |
| 1.67 | 0.791 | 0.951 | | 1.78 | k= 4, v=63 |
| 5 | 0.726 | 0.205 | | 1.79 | k= 5, v=63 |
| 15 | 0.477 | 2.646 | * | 1.79 | k= 6, v=63 |

s = 0.195

Note: df used for table values are approximate when v > 20.

Estimates of EC%

| Parameter | Estimate | 95% Bounds | | Std.Err. | Lower Bound /Estimate |
|-----------|----------|------------|-------|----------|--------------------------|
| | | Lower | Upper | | |
| EC5 | 2.9 | 0.74 | 11. | 0.29 | 0.26 |
| EC10 | 4.3 | 1.6 | 12. | 0.22 | 0.36 |
| EC25 | 8.7 | 5.2 | 15. | 0.11 | 0.59 |
| EC50 | 19. | 12. | 29. | 0.098 | 0.64 |

Slope = 2.01 Std.Err. = 0.840

Goodness of fit: p = 0.089 based on DF= 4.0 63.

5701WW : wheat dry weight

Observed vs. Predicted Treatment Group Means

| Dose | #Reps. | Obs. Mean | Pred. Mean | Obs. -Pred. | Pred. %Control | %Change |
|--------|--------|--------------|---------------|----------------|-------------------|----------|
| 0.00 | 10.0 | 0.708 | 0.826 | -0.118 | 100. | 0.00 |
| 0.0600 | 10.0 | 0.786 | 0.826 | -0.0395 | 100. | 2.53e-05 |
| 0.190 | 10.0 | 0.873 | 0.826 | 0.0471 | 100. | 0.00295 |
| 0.560 | 10.0 | 0.951 | 0.825 | 0.126 | 99.9 | 0.107 |
| 1.67 | 10.0 | 0.791 | 0.812 | -0.0210 | 98.3 | 1.72 |
| 5.00 | 10.0 | 0.726 | 0.724 | 0.00194 | 87.6 | 12.4 |
| 15.0 | 10.0 | 0.477 | 0.477 | 3.59e-05 | 57.8 | 42.2 |

!!!Warning: EC50 not bracketed by doses evaluated.

DP Barcode: D295614

MRID No.: 45745701

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